

09-965738substitute.ST25.txt SEQUENCE LISTING

```
O'Brien, Timothy J. Beard, John B. __
<110>
       Underwood, Lowell J.
       Repeat Sequences of the CA125 Gene and Their Use for Diagnostic
<120>
       and Therapeutic Interventions
       022438.43865
<130>
       us 09/965,738
<140>
       2001-09-27
<141>
<150>
       us 60/284,175
<151>
       2001-04-17
       us 60/299,380
<150>
<151>
       2001-06-19
<160>
       310
<170>
       PatentIn version 3.2
<210>
       13
<211>
<212>
       PRT
<213>
       Homo sapiens
<400>
Gln His Pro Gly Ser Arg Lys Phe Lys Thr Thr Glu Gly
<210>
<211>
       11
<212>
       PRT
<213>
      Homo sapiens
<400> 2
Phe Leu Thr Val Glu Arg Val Leu Gln Gly Leu
1 10
<210>
       8
<211>
<212>
       PRT
<213>
       Homo sapiens
<400>
      3
Asp Thr Tyr Val Gly Pro Leu Tyr
<210>
       4
<211>
       8
<212>
       PRT
<213>
       Homo sapiens
<400>
       4
```

Asp Gly Ala Ala Asn Gly Val Asp 1 5 <210> 240 <211> <212> DNA <213> Homo sapiens <220> <221> **CDS** (1)..(240)<222> <400> cgt cga cct ggc tct aga aag ttt aac acc acg gag aga gtc ctt cag Arg Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln 1 5 10 15 48 ggt ctg ctc agg cct gtg ttc aag aac acc agt gtt ggc cct ctg tac Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr 96 tct ggc tgc aga ctg acc ttg ctc agg ccc aag aag gat ggg gca gcc Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala 35 40 45 144 acc aaa gtg gat gcc atc tgc acc tac cgc cct gat ccc aaa agc cct Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro 50 60192 gga ctg gac aga gag cag cta tac tgg gag ctg agc cag ggt gat gca Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Gly Asp Ala 65 70 75 80 240 <210> 6 <211> 80 <212> PRT <213> Homo sapiens <400> Arg Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln
1 10 15 Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala 40 45 Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro 50 60 Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Gly Asp Ala 65 70 75 80

```
<210> 7
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Primer
<400> 7
                                                                                 20
ggagaggtt ctgcagggtc
<210> 8
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic Primer
<400> 8
Glu Arg Val Leu Gln Gly
<210>
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Primer
<400> 9
                                                                                 20
gtgaatggta tcaggagagg
<210> 10
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic Primer
<400> 10
Pro Leu Leu Ile Pro Phe
1 5
<210> 11
<211> 131
<212> PRT
<213> Homo sapiens
<400> 11
Glu Arg Val Leu Gln Gly Leu Leu Arg Ser Leu Phe Lys Ser Thr Ser 10 15
```

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30

Lys Asp Gly Thr Ala Thr Gly Val Asp Ala Ile Cys Thr His His Pro
35 40 45

Asp Pro Lys Ser Pro Arg Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ala Leu Asp 70 75 80

Asn Asp Ser Leu Phe Val Asn Gly Phe Thr His Arg Ser Ser Val Ser 85 90 95

Thr Thr Ser Thr Pro Gly Thr Pro Thr Val Tyr Leu Gly Ala Ser Lys 100 105 110

Thr Pro Ala Ser Ile Phe Gly Pro Ser Ala Ala Ser Pro Leu Leu Ile 115 120 125

Pro Phe Thr 130

<210> 12

<211> 130

<212> PRT <213> Homo sapiens

<400> 12

Glu Arg Val Leu Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser 10 15

Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Asp Gly Ala Ala Thr Arg Ala Asp Ala Val Cys Thr His Arg Pro 45

Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu 50 60

Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr 85 90 95

Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg

Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile

Pro Phe 130

<210> 13

<211> <212> 132

PRT

<213> Homo sapiens

<400>

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser 10 15

val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu 20 25 30

Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Ile His Arg Leu 35 40 45

Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu 50 55 60

Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly

Thr Pro Ser Ser Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu 115 120 125

Leu Ile Pro Phe 130

<210> 14

<211> 130

<212> PRT

<213> Homo sapiens

<400> 14

```
09-965738substitute.ST25.txt
Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser 1 10 15
Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30
Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu 35 40 45
    Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu 50 60
Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80
Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala
85 90 95
Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly
Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile
115 120 125
Pro Phe
    130
<210>
       15
<211>
      130
<212>
       PRT
<213>
       Homo sapiens
<400>
Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser
1 10 15
Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu
20 25 30
Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu 35 40 45
Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu 50 60
Ser Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80
```

```
09-965738substitute.ST25.txt
Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu
85 90 95
Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly 100 105 110
Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Phe Leu Ile
115 120 125
Pro Phe
     130
<210>
<211>
       130
<212>
        PRT
<213>
        Homo sapiens
<400>
        16
Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Ser Thr Ser 10 15
Ala Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu
20 25 30
Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu 35 40 45
Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60
Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80
Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro
85 90 95
Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly
Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile
115 120 125
Pro Phe
     130
<210>
        17
        130
<211>
```

<212>

<213>

PRT

Homo sapiens

<400> 17

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro 85 90 95

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly $100 \hspace{1cm} 105 \hspace{1cm} 110$

Thr Pro Ala Ser Leu Pro Gly His Ile Val Pro Gly Pro Leu Leu Ile 115 120 125

Pro Phe 130

<210> 18

<211> 131

<212> PRT

<213> Homo sapiens

<400> 18

Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val 35 40 45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Page 8 Arg Asp Ser Leu Tyr Val Asp Gly Phe Asn Pro Trp Ser Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly 100 105 110

Thr Pro Ser Pro Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile 115 120 125

Pro Phe Thr 130

65

<210> 19

<211> 131

<212> PRT

<213> Homo sapiens

<400> 19

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 70 75 80

Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro 85 90 95

Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu 100 105 110

Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Val 115 120 125

Pro Phe Thr 130

<210> 20

```
<211>
      130
```

<212> **PRT**

<213> Homo sapiens

<400> 20

Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser 1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu 20 25 30

Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 35 40 45

Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly

Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile 115 120 125

Pro Phe 130

<210>

21 131 <211>

<212> **PRT**

<213> Homo sapiens

<400> 21

Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser 1 10 15

Ile Gly Pro Leu Tyr Ser Ser Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys Thr His His Pro 35 40 45

Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu 50 60 Page 10

Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asp Gly Phe Thr His Trp Ser Pro Ile Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Ile Val Asn Leu Gly Thr Ser Gly 100 105 110

Ile Pro Pro Ser Leu Pro Glu Thr Thr Ala Thr Gly Pro Leu Leu Ile

Pro Phe Thr 130

<210> 22

282 <211>

PRT

<212> <213> Homo sapiens

<400> 22

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser 1 10 15

Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu 20 25 30

Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro 35 40 45

Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly

Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Page 11

Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu 145 150 160

Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190

Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220

His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu 225 230 235 240

Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr 245 250 255

Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser 260 265 270

Phe Pro Gly His Thr Glu Pro Gly Pro Leu 275 280

<400> 23

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser 1 10 15

Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu 20 25 30

Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro 35 40 45

Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80 Page 12

<210> 23

<211> 286

<212> PRT

<213> Homo sapiens

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu 85 90 95

Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly 100 105 110

Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn 130 135 140

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190

Ala Thr Arg Val Asp Ala Val Cys Thr Gln Arg Pro Asp Pro Lys Ser 195 200 205

Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr 210 215 220

His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu 225 230 235 240

Tyr Val Asn Gly Leu Thr His Gln Ser Ser Met Thr Thr Arg Thr 245 250 255

Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser 260 265 270

Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe 275 280 285

<210> 24

<211> 250

<212> PRT <213> Homo sapiens

<400> 24

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
1 5 10 15
Page 13

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
35 40 45 Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 70 75 80 Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Thr Ser Val Pro 85 90 95 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly 100 105 110 Thr Pro Phe Ser Leu Pro Ser Pro Ala Thr Ala Gly Pro Leu Leu Val 115 120 125 Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp 130 135 140 Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160 Gln Thr Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala 180 185 190 Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser 195 200 205 Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220 Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Trp Ile Pro

<210> 25

<211> 286

<212> PRT

<213> Homo sapiens

<400> 25

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu
35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 65 70 75 80

Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro 85 90 95

Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu 100 105 110

Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile 115 120 125

Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn 130 135 140

Met His His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu 145 150 160

Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala 180 185 190

Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln Ser 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr 210 215 220

Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Page 15 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser Thr

Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro 260 265 270

Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile Pro Phe 275 280 285

230

<210> 26

<211> 286

<212> PRT

<213> Homo sapiens

<400> 26

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 70 75 80

Arg Gly Ser Leu Tyr Val Asn Gly Phe Ser Arg Gln Ser Ser Met Thr 85 90 95

Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg

Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Page 16

Tyr Ser Gly Cys Arg Leu Thr Ser Leu Lys Pro Glu Lys Asp Gly Ala 180 185 190

Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220

His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240

Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr 245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser 260 265 270

Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile Pro Phe 275 280 285

<210> 27

<211> 286

<212> PRT

<213> Homo sapiens

<400> 27

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro 85 90 95

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Phe Gly Page 17 Thr Pro Ala Ser Leu His Gly His Thr Ala Pro Gly Pro Val Leu Val 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 130 135 140

Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala 180 185 190

Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220

Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu 225 230 235 240

Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr 245 250 255

Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser 260 265 270

Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile Pro Phe 275 280 285

<210> 28

<211> 286

<212> PRT

<213> Homo sapiens

<400> 28

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Page 18 Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60 Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro 85 90 95 Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly 100 105 110 Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val 115 120 125 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160 Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala 180 185 190 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220 Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr 245 250 255 Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser 260 265 270 Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile Pro Phe 275 280 285

<210> 29

<211> 281

<212> PRT

<213> Homo sapiens

<400> 29

Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val
35 40 45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly 100 105 110

Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn 130 135 140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala 180 185 190

Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser 195 200 205

Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220

```
O9-965738substitute.ST25.txt
Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu
225 230 235
Tyr Val Asn Gly Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr 245 250 255
Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu 260 265 270
Pro Ser Pro Thr Thr Ala Gly Pro Leu
<210>
<211>
        217
<212>
        PRT
        Homo sapiens
<213>
<400>
        30
Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser
1 10 15
Ile Gly Pro Leu Tyr Ser Ser Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30
Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys Thr His His Pro 45
Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu 50 60
Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80
Arg Asp Ser Leu Tyr Val Asp Gly Phe Thr His Trp Ser Pro Ile Pro 85 90 95
Thr Thr Ser Thr Pro Gly Thr Ser Ile Val Asn Leu Gly Thr Ser Gly
Ile Pro Pro Ser Leu Pro Glu Thr Thr Ala Thr Gly Pro Leu Leu Ile
Pro Phe Thr Pro Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp
Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu
```

09-965738substitute.ST25.txt Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr 210 215 31 286 <210> <211> <212> PRT Homo sapiens <213> <400> 31 Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser 1 10 15 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys 20 25 30 Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro
35 40 45 Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60 Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro 85 90 95 Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val 115 120 125 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu 145 150 155 160

09-965738substitute.ST25.txt Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220 His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Arg Thr 245 250 255 Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser 260 265 270Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe 275 280 285 <210> 32 <211> <212> 288 PRT <213> Homo sapiens <400> Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser 1 10 15 Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys
20 25 30 Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro $35 \ \ 40 \ \ 45$ Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60 Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro 85 90 95

09-965738substitute.ST25.txt Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu 145 150 155 160 Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala 180 185 190 Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser 195 200 205 Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220 Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr 245 250 255 Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser 260 265 270 Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe 275 <210> 33 <211> 284 <212> PRT <213> Homo sapiens <400> 33 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 1 5 10 15 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

09-965738substitute.ST25.txt Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 35 40 45 Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys Glu Leu 50 60 Ser Gln Leu Thr His Asp Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp 65 70 75 80 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro 85 90 95 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly 100 105 110 Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile 115 120 125 Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160 Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Glu Ala 180 185 190 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly 200 205 Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 220 Asn Ser Ile His Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr 245 250 255 Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser 260 265 270 Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile 275 280

<210> 34 288 <211> <212> PRT Homo sapiens <400> Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Ser Lys Asn Ser Ser 1 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu 20 25 30

Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 35 40 45

Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly $100 \hspace{1cm} 105 \hspace{1cm} 110$

Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile 115 120 125

Pro Phe Thr Val Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn 130 140

Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala 180 185 190

Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220

Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240 240 230

Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr 245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser 260 265 270

Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe 275 280 285

<210>

35 274 <211>

<212> PRT

<213> Homo sapiens

<400> 35

Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser 1 10 15

Val Gly Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro
35 40 45

Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu 50 60

Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80

Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr 85 90 95

Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg

Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val 115

Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr Glu Glu Asn 130 135 140 130

Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala 180 185 190

Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220

His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Gln Asp Arg Asp Ser Leu 235 235 240

Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile 245 250 255

Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser 260 265 270

Leu Pro

<210> 36

<211> 386 <212> PRT

<213> Homo sapiens

<400> 36

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser 1 10 15

Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 65 70 75 80

Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro 85 90 95

Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Ala Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160 Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190 Ala Thr Arg Val Asp Ala Ala Cys Thr Tyr Arg Pro Asp Pro Lys Ser 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220 His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Val Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr 245 250 255 Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser 260 265 270 Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu 275 280 285 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro 290 295 300 Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu 305 310 315 Arg Pro Leu Phe Lys Asn Thr Ser Ile Gly Pro Leu Tyr Ser Ser Cys 325 330 335 Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Lys Ala Ala Thr Arg Val 340 345 350 Page 29

Asp Ala Ile Cys Thr His His Pro Asp Pro Gln Ser Pro Gly Leu Asn 355 360 365

Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Thr 370 375 380

Glu Leu 385

<210> 37

<211> 438 <212> PRT

<212> PRT <213> Homo sapiens

<400> 37

Glu Arg Val Leu His Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Arg 1 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val 35 40 45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly 100 105 110

Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn 130 135 140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175 Page 30

Tyr Ser Gly Cys Arg Leu Thr Leu Phe Lys Pro Glu Lys His Glu Ala 180 185 190 Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly 195 200 205 Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 220 Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile 245 250 255 Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser 260 265 270 Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu 275 280 285 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr 290 295 300 Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu 305 310 315 320Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 325 330 335 Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala Ala Thr Gly Val 340 350 Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp 355 360 365 Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Arg Gly Ile Ile 370 375 380 Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu Tyr Val Asn Gly 385 390 395 400 Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Ile His Pro Ser Leu Pro Arg Pro Page 31

Ile Val Pro Gly Pro Leu 435

<210> 38

<211> 420 <212> PRT

<213> Homo sapiens

<400> 38

Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys $1 \hspace{1cm} 10 \hspace{1cm} 15$

Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn 20 25 30

Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser 35 40 45

Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg 50 60

Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr 65 70 75 80

Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr 85 90 95

Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro 100 105 110

Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met 115 120 125

Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Ser Val Leu Gln 130 135 140

Gly Leu Leu Thr Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr 145 150 155 160

Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala 165 170 175

Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln Ser Pro 180 185 190

Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr Asn Page 32 Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr 210 215 220

Val Asn Gly Phe Thr His Arg Ser Leu Gly Leu Thr Thr Ser Thr Pro 225 230 235 240

Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val 245 250 255

Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn 260 265 270

Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly 275 280 285

Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Arg 290 295 300

Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg 305 310 315 320

Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp 325 330 335

Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg 340 345 350

Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu 355 360 365

Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe 370 380

Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Pro Thr 385 390 395 400

Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser Lys Pro Gly Pro Ser 405 410 415

Ala Ala Ser Pro 420

<210> 39

<211> 439

<212> PRT <213> Homo sapiens

<400> 39

Glu Arg Val Leu Gln Gly Pro Leu Ser Pro Ile Phe Lys Asn Ser Ser 1 10 15

val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu 20 25 30

Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 35 40 45

Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly 100 105 110

Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn 130 135 140

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190

Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys Glu Leu Ser Gln Leu Thr 210 215 220

His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu 225 230 235 240

09-965738substitute.ST25.txt Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr 245 250 255 Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser 260 265 270 Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu 275 280 285 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu 310 Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 325 330 335 Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val 340 345 350 Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr 370 380 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu 435 <210> 40 <211> 424 <212> Homo sapiens <400> 40 Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg
1 10 15 09-965738substitute.ST25.txt
Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Tyr Thr His
20 25 30 Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp
35 40 45 Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 50 60 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser 65 70 75 80 Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr 85 90 95 Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu 100 Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg 130 135 140 Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly 145 150 155 160 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp 165 170 175 Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro 180 185 190 Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro Thr Thr 225 230 235 240 Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe 260

Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His 275 280 285

Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 290 295 300

Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser 305 310 315 320

Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr 325 330 335

Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly 340 345 350

Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser 360 365

Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val 370 380

Asn Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly 385 390 395

Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro 405 410 415

Gly His Thr Ala Pro Val Pro Leu 420

<210> 41

<211> 418

<212> PRT

<213> Homo sapiens

<400> 41

Thr Leu Leu Arg Pro Lys Lys Asp Gly Val Ala Thr Gly Val Asp Ala 1 5 10 15

Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu 20 25 30

Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu 35 40 45

Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr 50 60

His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Ser Thr Val 65 70 75 80 Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser Leu Ser Ser Pro Thr Ile 85 90 95 Met Ala Ala Gly Pro Leu Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile 100 105 110Thr Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys 115 120 125 Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu 145 150 155 160 Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys 165 170 175 Thr His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro 195 200 205 Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg 210 215 220 Ser Ser Met Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val 225 230 240 Gly Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly 245 250 255 Pro Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 315 Page 38

Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu 325 330 335

Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 340 345 350

Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp 355 360 365

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro 370 375 380

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly 385 390 395 400

Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile 405 410 415

Pro Phe

<210> 42

<211> 443

<212> PRT

<213> Homo sapiens

<400> 42

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Asp Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro 35 40 45

Asp Pro Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu 100 105 110 Page 39

Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Ser Glu 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Ser Gly Val Leu Cys Pro Pro Pro Ser Ile Leu Gly Ile Phe Thr Val Gln Pro Glu Thr Phe Glu Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 305 310 315 Leu Met Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 325 330 335 Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly 340 345 350 340 val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly Leu Page 40

Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile 370 375 380

Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn 385 390 395 400

Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr 405 410 415

Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly 420 425 430

His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe 435 440

<210> 43

<211> 442

<212> PRT

<213> Homo sapiens

<400> 43

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser 1 10 15

Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu 20 25 30

Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro 35 40 45

Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly 100 105 110

Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Page 41 Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu 145 150 155 160 Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala 180 185 190 Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser 195 200 205 Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220 Asn Asp Ile Glu Glu Val Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Arg Ser Phe Val Ala Pro Thr Ser Thr 245 250 255 Leu Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser 260 265 270 Leu Pro Ser Pro Thr Thr Gly Val Pro Leu Leu Ile Pro Phe Thr Leu 275 280 285 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro 290 295 300 Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu 305 310 315 320 Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys 325 330 335 Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val 340 345 350 Val Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp 355 360 365 Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr 370 375 380

09-965738substitute.ST25.txt Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe <210> 442 <211> <212> PRT Homo sapiens <213> <400> Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 65 70 75 80 Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro 85 90 95 Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu 145 150 155 160 09-965738substitute.ST25.txt Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu 165 170 175Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Ser Ser 180 185 190 Thr Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp 195 200 205 Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr 210 215 220 Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Arg Ser Phe Met Pro Thr Thr Ser Thr 245 250 255 Leu Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser 260 265 270 Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu 275 280 285 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr 290 295 300 Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu 305 310 315 320Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 325 330 335 Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly Val 340 345 350 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn 355 360 365 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu 370 375 380 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 385 390 395 Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Ser 410

Thr Val Asp Pro Arg Thr Ser Gly Thr Pro Ser Ser Leu Ser Ser Pro 420 425 430

Thr Ile Met Ala Ala Gly Pro Leu Leu Ile 435 440

<210> 45

<211> 379

<212> PRT <213> Homo sapiens

<400> 45

Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser 10 15

Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu 35 40 45

Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala 85 90 95

Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly 100 105 110

Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp 130 140

Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160

Gln Ser Leu Phe Gly Pro Met Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Phe Arg Ser Glu Lys Asp Gly Ala 180 185 190

Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser 195 200 205

Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220

Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240

Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro Asn Thr Ser Thr 245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser 260 265 270

Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu 275 280 285

Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr 290 295

Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu 305

Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 325 330 335

Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Val 340 345 350 340

Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn 360

Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu

<210> 46

<211> 439

<212> PRT

<213> Homo sapiens

<220>

MISC_FEATURE <221>

<222>

(1)..(439) Any "X" = any amino acid <223>

<400>

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Page 46

5

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu 35 40 45

Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro 85 90 95

Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly 100 105 110

Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn 130 140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu 145 150 155 160

Gln Gly Cys Leu Val Pro Cys Ser Arg Asn Thr Asn Val Gly Leu Leu 165 170 175

Xaa Xaa Xaa Xaa Xaa Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240

Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr 245 250 255

09-965738substitute.ST25.txt Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser 260 265 270 Leu Pro Ser Pro Thr Thr Val Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp 340 345 350 Ala Ile Cys Thr His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg 355 360 365 Glu Gln Leu Tyr Trp Gln Leu Ser Gln Val Thr Asn Gly Ile Lys Glu 370 375 380 Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser Thr Pro Trp Thr Ser Thr 405 410 415 Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val Pro Ser Pro Thr 420 425 430 Thr Ala Gly Pro Leu Leu Ile 435 <210> 47 <211> 1366 <212> PRT Homo sapiens <213> <400> 47 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser 1 10 15 Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu 20 25 30

09-965738substitute.ST25.txt Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro 35 40 45 Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60 Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro 85 90 95 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu 145 150 155 160 Gln Gly Pro Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220 His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu 225 Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr 245 250 255 Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser 260 265 270 Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu 275 280 285

Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu Gln Gly Leu Leu 305 310 315 Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys 325 330 335 Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Met 340 350 Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp 355 360 365 Arg Glu Gln Leu Tyr Cys Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His
420 425 430 Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe 465 470 475 480 Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu 485 490 495 Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys 500 510 Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu 515 520 525 Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro 535 540 Page 50

Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg 545 550 555 560 Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu 565 570 575 Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly 580 585 590 Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln 595 600 605 Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr 610 620 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 625 635 640 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 645 650 655 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 660 670 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 675 680 685 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 690 700 Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro 705 710 715 720 Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu 725 730 735 Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile 740 745 750 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn 755 760 765 Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu 770 775 780 Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu Page 51

Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser 810 815 Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 850 860 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu 930 Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 945 950 955 960 Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly Val 965 970 975 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn 980 985 990

Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu 995 1000 1005

Glu Val Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn 1010 1015 1020

Gly Phe Thr His Arg Ser Phe Val Ala Pro Thr Ser Thr Leu Gly 1025 1030 1035

09-965738substitute.ST25.txt Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu 1045 1040 Pro Ser Pro Thr Thr Gly Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly 1085 1090 1095 Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Ser Leu Tyr 1100 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 1115 1120 Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys 1130 1140 1130 Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly Ile Ile Glu Leu Gly Pro Tyr Thr Leu Asp Arg 1160 1165 1170His Ser Phe Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr 1230 1225 1220 Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr 1245 1235 Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr 1250 1260 Ser val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg 1265

Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr 1280 1285 1290

Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu 1295 1300 1305

Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly 1310 1315 1320

Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr 1325 1330 1335

His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala 1340 1345 1350

Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro 1355 1360 1365

<210> 48

<211> 1148

<212> PRT

<213> Homo sapiens

<400> 48

Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys 1 5 10 15

Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val 20 25 30

Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp 35 40 45

Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly Ile Ile 50 60

Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Phe Tyr Val Asn Gly 65 70 75 80

Phe Thr His Gln Ser Ser Met Thr Thr Thr Arg Thr Pro Asp Thr Ser 85 90 95

Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro
100 105 110

Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile 115 120 125

Thr Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe 145 150 155 160 Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu 165 170 175 Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys 180 185 190 Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu 195 200 205 Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro 210 215 220 Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg 225 230 235 240 Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu 245 250 255 Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly Pro Ser Ala Ala Ser 260 265 270 Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Arg 275 280 285 Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr 290 295 300 Glu Arg Val Leu Gln Gly Leu Leu Arg Ser Leu Phe Lys Ser Thr Ser 305 310 315 320 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 325 330 335 Lys Asp Gly Thr Ala Thr Gly Val Asp Ala Ile Cys Thr His His Pro 340 345 350 Asp Pro Lys Ser Pro Arg Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 355 360 365 Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly His Tyr Ala Leu Asp 370 380 Page 55

Asn Asp Ser Leu Phe Val Asn Gly Phe Thr His Arg Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Pro Thr Val Tyr Leu Gly Ala Ser Lys Thr Pro Ala Ser Ile Phe Gly Pro Ser Ala Ala Ser His Leu Leu Ile 420 425 430 Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn 435 440 445 Met Trp Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln 450 460 Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr 465 470 475 480 Ser Gly Ser Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Glu Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Gln Leu Tyr Leu Glu Leu Ser Gln Leu Thr His 515 520 525 Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr 530 540 Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Thr 545 555 val val Ser Glu Glu Pro Phe Thr Leu Asn Phe Thr Ile Asn Asn Leu Arg Tyr Met Ala Asp Met Gly Gln Pro Gly Ser Leu Lys Phe Asn Ile Thr Asp Asn Val Met Lys His Leu Leu Ser Pro Leu Phe Gln Arg Ser Ser Leu Gly Ala Arg Tyr Thr Gly Cys Arg Val Ile Ala Leu Arg Ser Val Lys Asn Gly Ala Glu Thr Arg Val Asp Leu Leu Cys Thr Tyr Leu Page 56

Gln Pro Leu Ser Gly Pro Gly Leu Pro Ile Lys Gln Val Phe His Glu 645 650 655 Leu Ser Gln Gln Thr His Gly Ile Thr Arg Leu Gly Pro Tyr Ser Leu 660 665 670 Asp Lys Asp Ser Leu Tyr Leu Asn Gly Tyr Asn Glu Pro Gly Leu Asp 675 680 685 Glu Pro Pro Thr Thr Pro Lys Pro Ala Thr Thr Phe Leu Pro Pro Leu 690 695 700 Ser Glu Ala Thr Thr Ala Met Gly Tyr His Leu Lys Thr Leu Thr Leu 705 710 715 720 Asn Phe Thr Ile Ser Asn Leu Gln Tyr Ser Pro Asp Met Gly Lys Gly 725 730 735 Ser Ala Thr Phe Asn Ser Thr Glu Gly Val Leu Gln His Leu Leu Arg 740 745 750 Pro Leu Phe Gln Lys Ser Ser Met Gly Pro Phe Tyr Leu Gly Cys Gln 755 760 765 Leu Ile Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp 770 775 780 Thr Thr Cys Thr Tyr His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile 785 790 795 800 Gln Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln 805 810 815 Leu Gly Phe Tyr Val Leu Asp Arg Asp Ser Leu Phe Ile Asn Gly Tyr 820 825 830 Ala Pro Gln Asn Leu Ser Ile Arg Gly Glu Tyr Gln Ile Asn Phe His 835 840 845 Ile Val Asn Trp Asn Leu Ser Asn Pro Asp Pro Thr Ser Ser Glu Tyr 850 860

Ile Thr Leu Leu Arg Asp Ile Gln Asp Lys Val Thr Thr Leu Tyr Lys 865 870 875 880

Gly	Ser	Gln	Leu	His 885	Asp	Thr					tute. Leu				Leu
Thr	Met	Asp	Ser 900	Val	Leu	۷a٦	Thr	va1 905	Lys	Ala	Leu	Phe	Ser 910	Ser	Asn
Leu	Asp	Pro 915	Ser	Leu	val	Glu	Gln 920	۷al	Phe	Leu	Asp	Lys 925	Thr	Leu	Asn
Ala	Ser 930	Phe	ніѕ	Trp	Leu	G]y 935	Ser	Thr	Tyr	Gln	Leu 940	val	Asp	Ile	His
Val 945	Thr	Glu	Met	Glu	Ser 950	Ser	٧a٦	туг	Gln	Pro 955	Thr	Ser	Ser	Ser	Ser 960
Thr	Gln	His	Phe	туг 965	Leu	Asn	Phe	Thr	Ile 970	Thr	Asn	Leu	Pro	Tyr 975	Ser
Gln	Asp	Lys	Ala 980	Gln	Pro	Gly	Thr	Thr 985	Asn	туг	Gln	Arg	Asn 990	Lys	Arg
Asn	Ile	G]u 995	Asp	Ala	Leu	Asn	Gln 100		u Pho	e Arq	g Asr	se 10		er I	le Lys
Ser	Tyr 1010		e Ser	^ Asp) Cys	G]r 101		al S	er Tl	nr Pł		-g 020	Ser '	val	Pro
Asn	Arg 1025		s His	s Thr	· Gly	Val 103		sp S	er Lo	eu Cy		sn)35	Phe :	Ser	Pro
Leu	Ala 1040		g Arg	y Val	Asp	Arc 104) V	al A	la I	le Ty	yr G ⁻ 10	lu 050	Glu	Phe	Leü
Arg	Met 1055		· Arg	g Asr	ı Gly	Thr 106		ln L∘	eu G	ln As	sn Pl 1(ne 065	Thr	Leu <i>i</i>	Asp
Arg	Ser 1070		· Va	l Leu	ı Val	Asp 107	o G	1у т	yr S	er Pi		sn 080	Arg /	Asn (Glu
Pro	Leu 1085		· Gly	/ Asr	ı Ser	Asp 109		eu P	ro Pl	he Ti		l a 095	val :	Ile	Leu
Ile		Lei	ı Ala	a Gly	/ Leu	Lei 110	ı G	ly L	eu I	le Ti	hr Cy 1.	/s 110	Leu :	Ile	Cys
Gly		Lei	ı Va ⁻	l Thr	Thr) A	rg A	rg L	ys L	ys G 1:	lu 125	Gly (Glu '	Tyr

Asn Val Gln Gln Gln Cys Pro Gly Tyr Tyr Gln Ser His Leu Asp 1130 1135 1140

Leu Glu Asp Leu Gln 1145

<210> 49 <211> 6833 <212> DNA

<213> Homo sapiens

<400> 49 gagagggttc tgcagggtct gctcaaaccc ttgttcagga atagcagtct ggaatacctc 60 tattcaggct gcagactagc ctcactcagg ccagagaagg atagctcagc catggcagtg 120 180 gatgccatct gcacacatcg ccctgaccct gaagacctcg gactggacag agagcgactg tactgggagc tgagcaatct gacaaatggc atccaggagc tgggccccta caccctggac 240 300 cggaacagtc tctatgtcaa tggtttcacc catcgaagct ctatgcccac caccagcact 360 cctqqqacct ccacaqtqqa tqtqqqaacc tcagggactc catcctccag ccccagcccc acgactgctg gccctctcct gatgccgttc accctcaact tcaccatcac caacctgcag 420 480 tacgaggagg acatgcgtcg cactggctcc aggaagttca acaccatgga gagggttctg cagggtccgc ttagtcccat attcaagaac tccagtgttg gccctctgta ctctggctgc 540 600 agactgacct ctctcaggcc cgagaaggat ggggcagcaa ctggaatgga tgctgtctgc 660 ctctaccacc ctaatcccaa aagacctggg ctggacagag agcagctgta ctgggagcta 720 agccagctga cccacaacat cactgagctg ggcccctaca gcctggacag ggacagtctc tatgtcaatg gtttcaccca tcagaactct gtgcccacca ccagtactcc tgggacctcc 780 acagtgtact gggcaaccac tgggactcca tcctccttcc ccggccacac agagcctggc 840 cctctcctga taccattcac gctcaacttc accatcacta acctacagta tgaggagaac 900 960 atgggtcacc ctggctccag gaagttcaac atcacggaga gggttctgca gggtctgctt 1020 aatcccattt tcaagaactc cagtgttggc cctctgtact ctggctgcag actgacctct 1080 ctcaggcccg agaaggatgg ggcagcaact ggaatggatg ctgtctgcct ctaccaccct 1140 aatcccaaaa qacctqqqct qqacaqaqaq cagctgtact gcgagctaag ccagctgacc cacaacatca ctgagctggg cccctacagc ttggacaggg acagtcttta tgtcaatggt 1200 ttcacccatc agaactctgt gcccaccacc agtactcctg ggacctccac agtgtactgg 1260 1320 gcaaccactg ggactccatc ctccttcccc ggccacacag agcctggccc tctcctgata 1380 ccattcaccc tcaacttcac catcaccaac ctgcagtacg aggaggacat gcgtcgcact ggctccagga agttcaacac catggagagg gttctgcagg gtctgctcaa gcccttgttc 1440

aagagcacca	gcgttggccc	tctgtactct	ggctgcagac		cagacctgag	1500
aaacatgggg	cagccactgg	agtggacgcc	atctgcaccc	tccgccttga	tcccactggt	1560
cctggactgg	acagagagcg	gctatactgg	gagctgagcc	agctgaccaa	cagcgttaca	1620
gagctgggcc	cctacaccct	ggacagggac	agtctctatg	tcaatggctt	cacccatcgg	1680
agctctgtgc	caaccaccag	tattcctggg	acctctgcag	tgcacctgga	aacctctggg	1740
actccagcct	ccctccctgg	ccacacagcc	cctggccctc	tcctggtgcc	attcaccctc	1800
aacttcacta	tcaccaacct	gcagtatgag	gaggacatgc	gtcaccctgg	ttccaggaag	1860
ttcaacacca	cggagagagt	cctgcagggt	ctgctcaagc	ccttgttcaa	gagcaccagt	1920
gttggccctc	tgtactctgg	ctgcagactg	accttgctca	ggcctgaaaa	acgtggggca	1980
gccaccggcg	tggacaccat	ctgcactcac	cgccttgacc	ctctaaaccc	tggactggac	2040
agagagcagc	tatactggga	gctgagcaaa	ctgacccgtg	gcatcatcga	gctgggcccc	2100
tacctcctgg	acagaggcag	tctctatgtc	aatggtttca	cccatcggaa	ctttgtgccc	2160
atcaccagca	ctcctgggac	ctccacagta	cacctaggaa	cctctgaaac	tccatcctcc	2220
ctacctagac	ccatagtgcc	tggccctctc	ctgataccat	tcacactcaa	cttcaccatc	2280
actaacctac	agtatgagga	gaacatgggt	caccctggct	ccaggaagtt	caacatcacg	2340
gagagggttc	tgcagggtct	gctcaaaccc	ttgttcagga	atagcagtct	ggaatacctc	2400
tattcaggct	gcagactaac	ctcactcagg	ccagagaagg	atagctcaac	catggcagtg	2460
gatgccatct	gcacacatcg	ccctgaccct	gaagacctcg	gactggacag	agagcgactg	2520
tactgggagc	tgagcaatct	gacaaatggc	atccaggagc	tgggccccta	caccctggac	2580
cggaacagtc	tctatgtcaa	tggtttcacc	catcgaagct	ctatgcccac	caccagcact	2640
cctgggacct	ccacagtgga	tgtgggaacc	tcagggactc	catcctccag	ccccagcccc	2700
acgactgctg	gccctctcct	gatgccgttc	accctcaact	tcaccatcac	caacctgcag	2760
tacgaggagg	acatgcgtcg	cactggctcc	aggaagttca	acaccatgga	gagtgtcctg	2820
cagggtctgc	tcaagccctt	gttcaagaac	accagtgttg	gccctctgta	ctctggctgc	2880
agattgacct	tgctcaggcc	caagaaagat	ggggcagcca	ctggagtgga	tgccatctgc	2940
acccaccgcc	ttgaccccaa	aagccctgga	ctcaacaggg	agcagctgta	ctgggagtta	3000
agcaaactga	ccaatgacat	tgaagaggtg	ggcccctaca	ccttggacag	gaacagtctc	3060
tatgtcaatg	gtttcaccca	tcggagcttt	gtggccccca	ccagcactct	tgggacctcc	3120
acagtggacc	ttgggacctc	agggactcca	tcctccctcc	ccagccccac	aacaggtgtt	3180
cctctcctga	taccattcac	actcaacttc	accatcacta	acctacagta	tgaggagaac	3240
atgggtcacc	ctggctccag	gaagttcaac	atcatggaga	gggttctgca	gggtctgctt	3300
atgcccttgt	tcaagaacac	cagtgtcagc	tctctgtact Page	ctggttgcag 60	actgaccttg	3360

ctcaggcctg	agaaggatgg	ggcagccacc	agagtggttg	ctgtctgcac	ccatcgtcct	3420
gaccccaaaa	gccctggact	ggacagagag	cggctgtact	ggaagctgag	ccagctgacc	3480
cacggcatca	ctgagctggg	cccctacacc	ctggacaggc	acagtctcta	tgtcaatggt	3540
ttcacccatc	agagctctat	gacgaccacc	agaactcctg	atacctccac	aatgcacctg	3600
gcaacctcga	gaactccagc	ctccctgtct	ggacctacga	ccgccagccc	tctcctgata	3660
ccattcacaa	ttaacttcac	catcactaac	ctgcggtatg	aggagaacat	gcatcaccct	3720
ggctctagaa	agtttaacac	cacggagaga	gtccttcagg	gtctgctcag	gcctgtgttc	3780
aagaacacca	gtgttggccc	tctgtactct	ggctgcagac	tgaccttgct	caggcccaag	3840
aaggatgggg	cagccaccaa	agtggatgcc	atctgcacct	accgccctga	tcccaaaagc	3900
cctggactgg	acagagagca	gctatactgg	gagctgagcc	agctaaccca	cagcatcact	3960
gagctgggcc	cctacaccct	ggacagggac	agtctctatg	tcaatggttt	cacacagcgg	4020
agctctgtgc	ccaccactag	cattcctggg	acccccacag	tggacctggg	aacatctggg	4080
actccagttt	ctaaacctgg	tccctcggct	gccagccctc	tcctggtgct	attcactctc	4140
aacttcacca	tcaccaacct	gcggtatgag	gagaacatgc	agcaccctgg	ctccaggaag	4200
ttcaacacca	cggagagggt	ccttcagggc	ctgctcaggt	ccctgttcaa	gagcaccagt	4260
gttggccctc	tgtactctgg	ctgcagactg	actttgctca	ggcctgaaaa	ggatgggaca	4320
gccactggag	tggatgccat	ctgcacccac	caccctgacc	ccaaaagccc	taggctggac	4380
agagagcagc	tgtattggga	gctgagccag	ctgacccaca	atatcactga	gctgggccac	4440
tatgccctgg	acaacgacag	cctctttgtc	aatggtttca	ctcatcggag	ctctgtgtcc	4500
accaccagca	ctcctgggac	ccccacagtg	tatctgggag	catctaagac	tccagcctcg	4560
atatttggcc	cttcagctgc	cagccatctc	ctgatactat	tcaccctcaa	cttcaccatc	4620
actaacctgc	ggtatgagga	gaacatgtgg	cctggctcca	ggaagttcaa	cactacagag	4680
agggtccttc	agggcctgct	aaggcccttg	ttcaagaaca	ccagtgttgg	ccctctgtac	4740
tctggctcca	ggctgacctt	gctcaggcca	gagaaagatg	gggaagccac	cggagtggat	4800
gccatctgca	cccaccgccc	tgaccccaca	ggccctgggc	tggacagaga	gcagctgtat	4860
ttggagctga	gccagctgac	ccacagcatc	actgagctgg	gcccctacac	actggacagg	4920
gacagtctct	atgtcaatgg	tttcacccat	cggagctctg	tacccaccac	cagcaccggg	4980
gtggtcagcg	aggagccatt	cacactgaac	ttcaccatca	acaacctgcg	ctacatggcg	5040
gacatgggcc	aacccggctc	cctcaagttc	aacatcacag	acaacgtcat	gaagcacctg	5100
ctcagtcctt	tgttccagag	gagcagcctg	ggtgcacggt	acacaggctg	cagggtcatc	5160
gcactaaggt	ctgtgaagaa	cggtgctgag	acacgggtgg	acctcctctg	cacctacctg	5220

09-965738substitute.ST25.txt 5280 cagcccctca gcggcccagg tctgcctatc aagcaggtgt tccatgagct gagccagcag 5340 acccatggca tcacccggct gggcccctac tctctggaca aagacagcct ctaccttaac 5400 ggttacaatg aacctggtct agatgagcct cctacaactc ccaagccagc caccacattc ctgcctcctc tgtcagaagc cacaacagcc atggggtacc acctgaagac cctcacactc 5460 5520 aacttcacca tctccaatct ccagtattca ccagatatgg gcaagggctc agctacattc aactccaccg agggggtcct tcagcacctg ctcagaccct tgttccagaa gagcagcatg 5580 ggccccttct acttgggttg ccaactgatc tccctcaggc ctgagaagga tggggcagcc 5640 5700 actggtgtgg acaccacctg cacctaccac cctgaccctg tgggccccgg gctggacata cagcagcttt actgggagct gagtcagctg acccatggtg tcacccaact gggcttctat 5760 gtcctggaca gggatagcct cttcatcaat ggctatgcac cccagaattt atcaatccgg 5820 5880 ggcgagtacc agataaattt ccacattgtc aactggaacc tcagtaatcc agaccccaca 5940 tcctcagagt acatcaccct gctgagggac atccaggaca aggtcaccac actctacaaa 6000 ggcagtcaac tacatgacac attccgcttc tgcctggtca ccaacttgac gatggactcc 6060 qtqttqqtca ctgtcaaggc attgttctcc tccaatttgg accccagcct ggtggagcaa 6120 gtctttctag ataagaccct gaatgcctca ttccattggc tgggctccac ctaccagttg gtggacatcc atgtgacaga aatggagtca tcagtttatc aaccaacaag cagctccagc 6180 6240 acccagcact tctacccgaa tttcaccatc accaacctac catattccca ggacaaagcc 6300 cagccaggca ccaccaatta ccagaggaac aaaaggaata ttgaggatgc gctcaaccaa 6360 ctcttccgaa acagcagcat caagagttat ttttctgact gtcaagtttc aacattcagg 6420 tctgtcccca acaggcacca caccggggtg gactccctgt gtaacttctc gccactggct cggagagtag acagagttgc catctatgag gaatttctgc ggatgacccg gaatggtacc 6480 6540 cagctgcaga acttcaccct ggacaggagc agtgtccttg tggatgggta ttctcccaac 6600 agaaatgagc ccttaactgg gaattctgac cttcccttct gggctgtcat cttcatcggc 6660 ttggcaggac tcctgggact catcacatgc ctgatctgcg gtgtcctggt gaccacccgc cggcggaaga aggaaggaga atacaacgtc cagcaacagt gcccaggcta ctaccagtca 6720 cacctagacc tggaggatct gcaatgactg gaacttgccg gtgcctgggg tgcctttccc 6780

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Page 62

ccagccaggg tccaaagaag cttggctggg gcagaaataa accatattgg tcg

6833

<210> 50 <211> 2248 <212> PRT

<213> Homo sapiens

<400> 50

5

Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu 20 25 30 Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro 35 40 45 Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60 Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro 85 90 95 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly 100 105 110 Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met 115 120 125 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu 145 150 155 160 Gln Gly Pro Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220 His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr 245 250 255

09-965738substitute.ST25.txt Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser 260 265 270 Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu 275 280 285 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu Gln Gly Leu Leu 305 310 315 Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys 325 330 335 Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Met 340 345 350 Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys Glu Leu Ser Gln Leu Thr His Asn Ile Thr 370 375 380 Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 385 395 400 Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe 465 Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys 500 510

Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu 515 520 525 Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro 530 540 Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg 545 550 555 560 Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu 565 570 575 Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly 580 585 590 Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr 610 615 620 Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 625 635 640 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 645 650 655 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 660 670 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 675 680 685 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 690 700 Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro 705 710 715 720 Ile Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu 725 730 735 Thr Pro Ser Ser Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile 740 745 750 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn 755 760 765 Page 65

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Arg Val Leu 770 780 Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu 785 790 795 800 800 Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser 805 810 815 Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp 830 Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr 840 Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 850 860 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr 865 Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 945 950 955 960 Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly Val 965 970 975 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn 980 985 990 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu 1000 995 Glu Val Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Page 66

Gly Phe Thr His Arg Ser Phe Val Ala Pro Thr Ser Thr Leu Gly 1030 1025 Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu 1040 1045 1050 1040 Pro Ser Pro Thr Thr Gly Val Pro Leu Leu Ile Pro Phe Thr Leu 1060 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His 1070 1075 1080 Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Ser Leu Tyr 1100 1110 1110Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 1115 1120 1125 Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys 1140 1135 1130 Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln 1145 1150 1155 Leu Thr His Gly Ile Ile Glu Leu Gly Pro Tyr Thr Leu Asp Arg 1160 His Ser Phe Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr 1175 Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser 1190 Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu 1205 1210 1215 Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr 1240

09-965738substitute.ST25.txt Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu 1295 1300 1305 Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly 1320 1310 Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala 1345 1340 val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr 1370 1375 1380 Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 1420 Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Thr Ala Thr Gly 1430 1435 val Asp Ala Ile Cys Thr His His Pro Asp Pro Lys Ser Pro Arg 1445 1450 Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His 1470 1460 Asn Ile Thr Glu Leu Gly His Tyr Ala Leu Asp Asn Asp Ser Leu 1485 1475

Phe Val Asn Gly Phe Thr His Arg Ser Ser Val Ser Thr Thr Ser 1490 1500 Thr Pro Gly Thr Pro Thr Val Tyr Leu Gly Ala Ser Lys Thr Pro 1505 Ala Ser Ile Phe Gly Pro Ser Ala Ala Ser His Leu Leu Ile Leu 1520 1530 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn 1535 1540 1545 Met Trp Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 1550 1560 Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Ser Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Glu Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Gln Leu Tyr Leu Glu Leu 1610 1620 Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu 1625 Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser 1640 Val Pro Thr Thr Ser Thr Gly Val Val Ser Glu Glu Pro Phe Thr 1655 Leu Asn Phe Thr Ile Asn Asn Leu Arg Tyr Met Ala Asp Met Gly 1670 1675 Gln Pro Gly Ser Leu Lys Phe Asn Ile Thr Asp Asn Val Met Lys 1690 1685 His Leu Leu Ser Pro Leu Phe Gln Arg Ser Ser Leu Gly Ala Arg 1700 Tyr Thr Gly Cys Arg Val Ile Ala Leu Arg Ser Val Lys Asn Gly 1720 1725 Page 69

Page 70

Ala Glu Thr Arg Val Asp Leu Leu Cys Thr Tyr Leu Gln Pro Leu Ser Gly Pro Gly Leu Pro Ile Lys Gln Val Phe His Glu Leu Ser Gln Gln Thr His Gly Ile Thr Arg Leu Gly Pro Tyr Ser Leu Asp 1760 1765 1770 Lys Asp Ser Leu Tyr Leu Asn Gly Tyr Asn Glu Pro Gly Leu Asp 1775 1780 1785 Glu Pro Pro Thr Thr Pro Lys Pro Ala Thr Thr Phe Leu Pro Pro 1790 1795 1800 Leu Ser Glu Ala Thr Thr Ala Met Gly Tyr His Leu Lys Thr Leu 1805 Thr Leu Asn Phe Thr Ile Ser Asn Leu Gln Tyr Ser Pro Asp Met 1820 Gly Lys Gly Ser Ala Thr Phe Asn Ser Thr Glu Gly Val Leu Gln 1840 His Leu Leu Arg Pro Leu Phe Gln Lys Ser Ser Met Gly Pro Phe 1855 1850 Tyr Leu Gly Cys Gln Leu Ile Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr Thr Cys Thr Tyr His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile Gln Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln Leu Gly Phe Tyr Val Leu Asp 1920 1910 1915 Arg Asp Ser Leu Phe Ile Asn Gly Tyr Ala Pro Gln Asn Leu Ser 1925 Ile Arg Gly Glu Tyr Gln Ile Asn Phe His Ile Val Asn Trp Asn 1945 1950 Leu Ser Asn Pro Asp Pro Thr Ser Ser Glu Tyr Ile Thr Leu Leu Arg Asp Ile Gln Asp Lys Val Thr Thr Leu Tyr Lys Gly Ser Gln Leu His Asp Thr Phe Arg Phe Cys Leu Val Thr Asn Leu Thr Met 1985 1990 1995 Asp Ser Val Leu Val Thr Val Lys Ala Leu Phe Ser Ser Asn Leu Asp Pro Ser Leu Val Glu Gln Val Phe Leu Asp Lys Thr Leu Asn 2015 2020 2025 Ala Ser Phe His Trp Leu Gly Ser Thr Tyr Gln Leu Val Asp Ile 2030 2040 His Val Thr Glu Met Glu Ser Ser Val Tyr Gln Pro Thr Ser Ser Ser Ser Thr Gln His Phe Tyr Leu Asn Phe Thr Ile Thr Asn Leu Pro Tyr Ser Gln Asp Lys Ala Gln Pro Gly Thr Thr Asn Tyr Gln 2080 Arg Asn Lys Arg Asn Ile Glu Asp Ala Leu Asn Gln Leu Phe Arg 2090 2100 Asn Ser Ser Ile Lys Ser Tyr Phe Ser Asp Cys Gln Val Ser Thr 2115 2105 Phe Arg Ser Val Pro Asn Arg His His Thr Gly Val Asp Ser Leu 2120 2130 21Ž0 Cys Asn Phe Ser Pro Leu Ala Arg Arg Val Asp Arg Val Ala Ile 2135 2140 2145 Tyr Glu Glu Phe Leu Arg Met Thr Arg Asn Gly Thr Gln Leu Gln 2150 2160 2160 Asn Phe Thr Leu Asp Arg Ser Ser Val Leu Val Asp Gly Tyr Ser 2165 2175 Pro Asn Arg Asn Glu Pro Leu Thr Gly Asn Ser Asp Leu Pro Phe

					09	9-965	57389	subst	ti tu1	te.ST	25.tx	κt		
Trp Ala 219		Ile	Leu	Ile	Gly 2200	Leu	Ala	Gly	Leu	Leu 2205	Gly	Leu	Ile	
Thr Cy:	s Leu 10	Ile	Cys	Gly	val 2215	Leu	Val	Thr	Thr	Arg 2220	Arg	Arg	Lys	
Lys Gl		Glu	Tyr	Asn	Val 2230	Gln	Gln	Gln	Cys	Pro 2235	Gly	Tyr	Tyr	
Gln Se 22		Leu	Asp	Leu	Glu 2245	Asp	Leu	Gln						
<210> <211> <212> <213>		icia	l Sed	quen	ce									
<220> <223>	Synth	etic	Prin	ner										
<400> cagcag	51 agac ca	agca	cgagt	t ac	tc									24
<210> <211> <212> <213>		icia ⁻	l Sed	quen	ce									
<220> <223>	Synth	etic	Prin	ner										
<400> tccact	52 gcca t	ggcto	gagci	t										20
<210> <211> <212> <213>	53 22 DNA Artif	icia ⁻	l Sed	quen	ce									
<220> <223>	Synth	etic	Pri	mer										
<400> ccagca	53 cagc t	cttc	ccag	g ac										22
<210> <211> <212> <213>	54 22 DNA Artif	icia ⁻	1 Se	quen	ce									
<220> <223>	Synth	etic	Pri	mer										
<400> ggaatg	54 gctg a	gctga	acgt	c tg										22

<210> <211> <212> <213>	55 21 DNA Artificial Sequence	
<220> <223>	Synthetic Primer	
<400> cttccc	55 agga caacctcaag g	21
<210> <211> <212> <213>	56 21 DNA Artificial Sequence	
<220> <223>	Synthetic Primer	
<400> gcagga	56 tgag tgagccacgt g	21
<210> <211> <212> <213>		
<220> <223>	Synthetic Primer	
<400> gtcaga	57 tctg gtgacctcac tg	22
<210> <211> <212> <213>	58 21 DNA Artificial Sequence	
<220> <223>	Synthetic Primer	
<400> gaggca	58 ctgg aaagcccaga g	21
<210> <211> <212> <213>	59 25 DNA Artificial Sequence	
<220> <223>	Synthetic Primer	
<400> ctgatg	59 gcat tatggaacac atcac	25
<210> <211>	60 22	

	09-965738substitute.ST25.txt	
<212> <213>	DNA Artificial Sequence	
<220> <223>	Synthetic Primer	
<400> cccagaa	60 acga gagaccagtg ag	22
<210> <211> <212> <213>		
<220> <223>	Synthetic Primer	
<400> gctgate	61 ggcg atgaatgaac actg	24
<210> <211> <212> <213>	22	
<220> <223>	Synthetic Primer	
<400> cccaga	62 acga gagaccagtg ag	22
<210> <211> <212> <213>	35	
<220> <223>	Synthetic Primer	
<400> cgcgga	63 tccg aacactgcgt ttgctggctt tgatg	35
	64 23 DNA Artificial Sequence	
<220> <223>	Synthetic Primer	
<400> cctctg	64 tgtg ctgcttcatt ggg	23
<210><211><211><212>	65 32 DNA Artificial Sequence	

09-965738substitute.ST25.txt <220> <223> Synthetic Primer <400> 65 32 accggatcca tgggccacac agagcctggc cc <210> 66 <211> 29 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Primer <400> 66 29 tgtaagctta ggcagggagg atggagtcc <210> 67 507 <211> <212> DNA <213> Homo sapien <400> atgagaggat cgcatcacca tcaccatcac ggatccatgg gccacacaga gcctggccct 60 120 ctcctgatac cattcacttt caactttacc atcaccaacc tgcattatga ggaaaacatg caacacctg gttccaggaa gttcaacacc acggagaggg ttctgcaggg tctgctcaag 180 cccttgttca agaacaccag tgttggccct ctgtactctg gctgcagact gaccttgctc 240 agacctgaga agcatgaggc agccactgga gtggacacca tctgtaccca ccgcgttgat 300 360 cccatcggac ctggactgga cagagagcgg ctatactggg agctgagcca gctgaccaac agcatcacag agctgggacc ctacaccctg gacagggaca gtctctatgt caatggcttc 420 480 aaccctcgga gctctgtgcc aaccaccagc actcctggga cctccacagt gcacctggca 507 acctctqqqa ctccatcctc cctgcct <210> 68 <211> 169 <212> PRT Homo sapiens <213> <400> 68 Met Arg Gly Ser His His His His His Gly Ser Met Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe

```
09-965738substitute.ST25.txt
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys
Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu 65 70 75 80
Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr
85 90 95
His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr
100 105 110
Tṛp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr
115 120 125
Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser
Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala
145 150 155 160
Thr Ser Gly Thr Pro Ser Ser Leu Pro
<210>
        69
<211>
        909
<212>
        PRT
<213> Homo sapiens
<220>
<221> MISC_FEATURE
<222> (1)..(909)
<223> Any "X" = any amino acid
<400>
Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser 10 15
Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys
20 25 30
Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro 35 40 45
Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60
Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 __ 80
                                            Page 76
```

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro 85 90 95 Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly 100 105 110 Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Ile 115 120 125 Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn 130 135 140 Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu 145 150 155 160 Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala 180 185 190 Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser 195 200 205 Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220 Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr 245 250 255 Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser 260 265 270 Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe 275 280 285 Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly 305 310 315 Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Page 77

Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr 340 345 350 340 Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser Pro Gly 355 360 365 Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly 370 380 Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val 385 390 400 Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr Pro Gly 405 410 415 Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser Pro 420 425 430 Ser Pro Thr Thr Ala Gly Pro Leu Leu Met Pro Phe Thr Leu Asn Phe 435 440 445 Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser 450 460 Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys Pro 465 470 475 480 Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu
485
490 Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala 500 505 510 Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu 515 520 525 Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu Leu 530 540 Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr 545 550 555 560 His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val 565 570 575

09-965738substitute.ST25.txt His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr Ala 580 585 590 Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Cys Leu Val Pro Cys Ser Arg Asn Thr Asn Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg 645 650 655 Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa 660 665 670 Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp 675 680 685 Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Thr 690 695 700 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser 705 715 720 Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr 725 730 735 Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Val Pro Leu Leu 740 745 750 Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu 755 760 765 Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 770 780 Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Gly Pro 785 790 795 800 Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly 805 810 815 Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln 820 830

Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln Val 835 840 845

Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser 850 860

Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser 865 870 875 880

Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser 885 890 895

Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile 900 905

<210> 70

<211> 525

<212> PRT

<213> Homo sapiens

<400> 70

Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu 1 5 10 15

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala 20 25 30

Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn 35 40 45

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 50 60

Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu 65 70 75 80

Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr 85 90 95

Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser 100 105 110

Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Val Pro Phe Thr Leu 115 120 125

Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Ala Met Arg His Pro 130 135 140

Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu 145 150 155 160 Arg Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys 165 170 175 Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val 180 185 190 Asp Ala Ala Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp 195 200 205 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr 210 215 220 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Val Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser 245 250 255 Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile 280 Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys 290 295 Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe 305 310 315 320 Lys Asn Thr Ser Ile Gly Pro Leu Tyr Ser Ser Cys Arg Leu Thr Leu 325 330 335 Leu Arg Pro Glu Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys 340 345 350 Thr His His Pro Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu 355 360 365 Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro 370 375 380 Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asp Gly Phe Thr His Trp 385 390 395 400 Page 81

Ser Pro Ile Pro Thr Thr Ser Thr Pro Gly Thr Ser Ile Val Asn Leu 405 410 415

Gly Thr Ser Gly Ile Pro Pro Ser Leu Pro Glu Thr Thr Ala Thr Gly 420 425 430

Pro Leu Leu Ile Pro Phe Thr Pro Asn Phe Thr Ile Thr Asn Leu Gln 435 440 445

Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met 450 460

Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser 465 470 475 480

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu 485 490 495

Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 500 510

Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr 515 520 525

<210> 71

<211> 594 <212> PRT

<213> Homo sapiens

<400> 71

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Asp Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro
35 40 45

Asp Pro Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro 85 90 95 Page 82

Thr Thr Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu 115 120 125 Phe Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 160 150 Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala 180 185 190 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Ser Glu 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Ser Gly Val Leu Cys Pro Pro Pro Ser Ile Leu Gly Ile Phe Thr Val Gln Pro Glu Thr Phe Glu Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His Arg 300 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 305 310 315 Leu Thr Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 325 330 335 Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly Page 83

Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly Leu 355 360 365 Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile 370 375 380 Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn 385 390 395 400 Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr $405 \hspace{1cm} 410 \hspace{1cm} 415$ Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly 420 430 His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr 435 440 445 Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg 450 455 460 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu 465 470 475 480 Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr 485 490 495 Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile 500 505 510 500 Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln 515 520 525 Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly 530 540 Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His 545 550 555 560 Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr Val Asp 565 570 575

Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Gly 580 585 590

72 424 <210> <211> <212> <213> **PRT** Homo sapiens <400> 72 Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Tyr Thr His $20 \hspace{1cm} 25 \hspace{1cm} 30$ Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp 35 40 45 Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 50 60 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser 65 70 75 80 Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr 85 90 95 Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu 100 105 110 Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Met Glu Arg 130 140Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly 145 150 155 160 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp 165 170 175 Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro 180 185 190 Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu Ser Gln 195 200 205

09-965738substitute.ST25.txt Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp 210 215 220 Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro Thr Thr 225 230 235 240 Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Ile Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 290 295 300 Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly 340 345 350 Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser 355 Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val 370 375 380 Asn Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly 385 390 395 400 Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu 420 <210> 73 <211> 286 <212> PRT <213> Homo sapiens

<400>

73

09-965738substitute.ST25.txt
Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser
1 10 15 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30 Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45 Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 55 60 Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 65 70 75 80 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro 85 90 95 Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly 100 105 110 Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val 115 120 125 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 130 135 140 Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160 Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu 165 170 175 Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala 180 185 190 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220 Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu 225 230 235 240 Tyr Val Asn Gly Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr 245 250 255

Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser 260 265 270

Leu Pro Arg Pro Ile Val Pro Gly Pro Leu Leu Ile Pro Phe 275 280 285

<210> 74

<211> 286

<212> PRT

<213> Homo sapiens

<400> 74

Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser 1 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys 20 25 30

Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro 35 40 45

Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro 85 90 95

Thr Thr Ser Ile Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly 100 105 110

Thr Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 130 135 140

Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190

Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 220

His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu 225 230 235 240

Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Arg Thr 245 250 255

Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser 260 265 270

Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe 275 280 285

<210> 75

<211> 286

<212> PRT

<213> Homo sapiens

<400> 75

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 35 40 45

Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60

Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp 65 70 75 80

Arg Gly Ser Leu Tyr Val Asn Gly Phe Ser Arg Gln Ser Ser Met Thr 85 90 95

Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg 100 105 110

Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn 130 135 140

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Asn Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Ser Leu Lys Pro Glu Lys Asp Gly Ala 180 185 190

Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg 195 200 205

Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr 210 215 220

His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 230 235 240

Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr 245 250 255

Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser 260 265 270

Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Ile Pro Phe 275 280 285

<210> 76

<211> 286

<212> PRT

<213> Homo sapiens

<400> 76

Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser 1 10 15

Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu 20 25 30

Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro 35 40 . 45

Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp 65 70 75 80

Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu 85 90 95

Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly 100 105 110

Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn 130 135 140

Met Gly His Pro Gly Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 180 185 190

Ala Thr Arg Val Asp Ala Val Cys Thr Gln Arg Pro Asp Pro Lys Ser 195 200 205

Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr 210 220

His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu 225 230 235 240

Tyr Val Asn Gly Leu Thr His Gln Ser Ser Met Thr Thr Arg Thr 245 250 255

Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser 260 265 270

Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Ile Pro Phe 275 280 285

<210> 77

²⁸⁸

<211> <212> <213> PRT

Homo sapiens

<400> 77

Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Ser Lys Asn Ser Ser 1 10 15 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu 20 25 30 Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 35 40 45 Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 50 60 Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp 65 70 75 80 Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro 85 90 95 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly 100 105 110 Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Val Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn 130 140 Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160 Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175 Tyr Ser, Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala 180 185 190 Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser 195 200 205 Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr 210 215 220 Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 225 235 240 Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr 245 250 255 Page 92

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser 260 265 270

Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu Ile Pro Phe 275 280 285

<210> 78

<211> 597

<212> PRT <213> Homo sapiens

<400> 78

Glu Arg Val Leu His Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Arg 1 5 10 15

Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu 20 25 30

Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val
35 40 45

Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 50 60

Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp 70 75 80

Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro 85 90 95

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly
100 105 110

Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile 115 120 125

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn 130 135 140

Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 145 150 155 160

Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu 165 170 175

Tyr Ser Gly Cys Arg Leu Thr Leu Phe Lys Pro Glu Lys His Glu Ala 180 185 190 Page 93

Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly 200 205 Pro Gly Leu Asp Arg Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn 210 220 Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr 225 230 235 240 Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro 245 250 255 Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu 260 265 270 Pro Gly His Thr Ala Pro Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn 275 280 285 Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly 290 295 300 Ser Arg Lys Phe Asn Thr Met Glu Arg Val Leu Gln Gly Leu Leu Lys 305 310 315 320Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg 325 330 335 Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala Ala Thr Gly Val Asp $340 \hspace{1cm} 345 \hspace{1cm} 350$ Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp Arg 355 360 365 Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Arg Gly Ile Ile Glu 370 380 Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu Tyr Val Asn Gly Phe 385 390 395 400 Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr Pro Gly Thr Ser Thr 405 410 415 Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser Leu Pro Arg Pro Ile 420 425 430 Val Pro Gly Pro Leu Leu Ile Pro Phe Thr Ile Asn Phe Thr Ile Thr Page 94

Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe 450 455 460

Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys 465 470 475 480

Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu 485 490 495

Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr 500 510

His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr 515 520 525

Trp Gln Leu Ser Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr 530 535 540

Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser 545 550 555 560

Ser Gly Leu Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly 565 570 575

Thr Ser Gly Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro 580 585 590

Leu Leu Ile Pro Phe 595

<210> 79

<211> 420

<212> PRT

<213> Homo sapiens

<400> 79

Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys
1 10 15

Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn 20 25 30

Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser 35 40 45

Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Page 95 Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr 65 70 75 80 Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr 85 90 95 Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro 100 105 110 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met 115 120 125 Gly His Pro Gly Ser Arg Lys Phe Asn Ile Thr Glu Ser Val Leu Gln 130 140 Gly Leu Leu Thr Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr 145 150 155 160 Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala 165 170 175 Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln Ser Pro 180 185 190 Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr Asn 195 200 205 Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr 210 215 220 Val Asn Gly Phe Thr His Arg Ser Leu Gly Leu Thr Thr Ser Thr Pro 225 230 235 240 Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val 245 250 255 Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn 260 265 270 Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly 275 280 285 Ser Arg Lys Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Arg 290 295 300

```
09-965738substitute.ST25.txt
Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg
Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp 325 330 335
Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg
Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu
Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe 370 380
Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Pro Thr
                       390
Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser Lys Pro Gly Pro Ser
Ala Ala Ser Pro
<210>
        80
        479
<211>
        PRT
<213>
        Homo sapiens
<400>
Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr 20^{\circ} 25^{\circ} 30
His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Ser Thr Val 35 40 45
Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser Leu Ser Ser Pro Thr Ile 50 60
Met Ala Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile 65 70 75 80
Thr Asn Leu Gln Tyr Glu Glu Asn Met Gly His Pro Gly Ser Arg Lys 85 90 95
```

09-965738substitute.ST25.txt Phe Asn Ile Met Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe 100 105 110 Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu 115 120 125 Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys 130 135 140 Ser His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu 145 150 155 160 Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro 165 170 175 Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg 180 185 190 Ser Ser Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val 210 215 220 Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys 235 230 240 Tyr Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr 245 250 255 Glu Arg Val Leu Gln Ser Leu Phe Gly Pro Met Phe Lys Asn Thr Ser 260 265 270Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu 275 280 285 Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu 290 295 300 Asp Pro Lys Ser Leu Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu 305 310 315 320 Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp 325 330 335 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro

Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly 355 360 365

Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val 370 375 380

Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 385 390 395 400

Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu 405 410 415

Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu 420 425 430

Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 435 440 445

Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser 450 455 460

Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu 465 470 475

<210> 81 <211> 5465 <212> DNA

<213> Homo sapiens

<400> 81
cagagagcgt tgagctggga acagtgacaa gtgcttatca agttccttca ctctcaacac 60
ggttgacaag aactgatggc attatggaac acatcacaaa aatacccaat gaagcagcac 120
acagaggtac cataagacca gtcaaaggcc ctcagacatc cacttcgcct gccagtccta 180

aaggactaca cacaggaggg acaaaaagaa tggagaccac caccacagct ttgaagacca 240

ccaccacage tttgaagace acttecagag ccacettgae caccagtgte tatactecca 300

ctttgggaac actgactccc ctcaatgcat caaggcaaat ggccagcaca atcctcacag 360

aaatgatgat cacaacccca tatgttttcc ctgatgttcc agaaacgaca tcctcattgg 420

ctaccagcct gggagcagaa accagcacag ctcttcccag gacaacccca tctgttctca 480

atagagaatc agagaccaca gcctcactgg tctctcgttc tggggcagag agaagtccgg 540

ttattcaaac tctagatgtt tcttctagtg agccagatac aacagcttca tgggttatcc 600

atcctgcaga gaccatccca actgtttcca agacaacccc caatttttc cacagtgaat 660

tagacactgt atcttccaca gccaccagtc atggggcaga cgtcagctca gccattccaa 720

caaatatctc	acctagtgaa			ggtcactatt		780
atactagtac	aacattccca	acactgacta	agtccccaca	tgaaacagag	acaagaacca	840
catggctcac	tcatcctgca	gagaccagct	caactattcc	cagaacaatc	cccaatttt	900
ctcatcatga	atcagatgcc	acaccttcaa	tagccaccag	tcctggggca	gaaaccagtt	960
cagctattcc	aattatgact	gtctcacctg	gtgcagaaga	tctggtgacc	tcacaggtca	1020
ctagttctgg	gacagacaga	aatatgacta	ttccaacttt	gactctttct	cctggtgaac	1080
caaagacgat	agcctcatta	gtcacccatc	ctgaagcaca	gacaagttcg	gccattccaa	1140
cttcaactat	ctcgcctgct	gtatcacggt	tggtgacctc	aatggtcacc	agtttggcgg	1200
caaagacaag	tacaactaat	cgagctctga	caaactcccc	tggtgaacca	gctacaacag	1260
tttcattggt	cacgcatcct	gcacagacca	gcccaacagt	tccctggaca	acttccattt	1320
ttttccatag	taaatcagac	accacacctt	caatgaccac	cagtcatggg	gcagaatcca	1380
gttcagctgt	tccaactcca	actgtttcaa	ctgaggtacc	aggagtagtg	acccctttgg	1440
tcaccagttc	tagggcagtg	atcagtacaa	ctattccaat	tctgactctt	tctcctggtg	1500
aaccagagac	cacaccttca	atggccacca	gtcatgggga	agaagccagt	tctgctattc	1560
caactccaac	tgtttcacct	ggggtaccag	gagtggtgac	ctctctggtc	actagttcta	1620
gggcagtgac	tagtacaact	attccaattc	tgactttttc	tcttggtgaa	ccagagacca	1680
caccttcaat	ggccaccagt	catgggacag	aagctggctc	agctgttcca	actgttttac	1740
ctgaggtacc	aggaatggtg	acctctctgg	ttgctagttc	tagggcagta	accagtacaa	1800
ctcttccaac	tctgactctt	tctcctggtg	aaccagagac	cacaccttca	atggccacca	1860
gtcatggggc	agaagccagc	tcaactgttc	caactgtttc	acctgaggta	ccaggagtgg	1920
tgacctctct	ggtcactagt	tctagtggag	taaacagtac	aagtattcca	actctgattc	1980
tttctcctgg	tgaactagaa	accacacctt	caatggccac	cagtcatggg	gcagaagcca	2040
gctcagctgt	tccaactcca	actgtttcac	ctggggtatc	aggagtggtg	acccctctgg	2100
tcactagttc	cagggcagtg	accagtacaa	ctattccaat	tctaactctt	tcttctagtg	2160
agccagagac	cacaccttca	atggccacca	gtcatggggt	agaagccagc	tcagctgttc	2220
taactgtttc	acctgaggta	ccaggaatgg	tgacctctct	ggtcactagt	tctagagcag	2280
taaccagtac	aactattcca	actctgacta	tttcttctga	tgaaccagag	accacaactt	2340
cattggtcac	ccattctgag	gcaaagatga	tttcagccat	tccaacttta	gctgtctccc	2400
ctactgtaca	agggctggtg	acttcactgg	tcactagttc	tgggtcagag	accagtgcgt	2460
tttcaaatct	aactgttgcc	tcaagtcaac	cagagaccat	agactcatgg	gtcgctcatc	2520
ctgggacaga	agcaagttct	gttgttccaa	ctttgactgt	ctccactggt	gagccgttta	2580
caaatatctc	attggtcacc	catcctgcag	agagtagctc Page	aactcttccc 100	aggacaacct	2640

caaggttttc	ccacagtgaa	ttagacacta	tgccttctac	agtcaccagt	cctgaggcag	2700
aatccagctc	agccatttca	actactattt	cacctggtat	accaggtgtg	ctgacatcac	2760
tggtcactag	ctctgggaga	gacatcagtg	caacttttcc	aacagtgcct	gagtccccac	2820
atgaatcaga	ggcaacagcc	tcatgggtta	ctcatcctgc	agtcaccagc	acaacagttc	2880
ccaggacaac	ccctaattat	tctcatagtg	aaccagacac	cacaccatca	atagccacca	2940
gtcctggggc	agaagccact	tcagattttc	caacaataac	tgtctcacct	gatgtaccag	3000
atatggtaac	ctcacaggtc	actagttctg	ggacagacac	cagtataact	attccaactc	3060
tgactctttc	ttctggtgag	ccagagacca	caacctcatt	tatcacctat	tctgagacac	3120
acacaagttc	agccattcca	actctccctg	tctccctgg	tgcatcaaag	atgctgacct	3180
cactggtcat	cagttctggg	acagacagca	ctacaacttt	cccaacactg	acggagaccc	3240
catatgaacc	agagacaaca	gccatacagc	tcattcatcc	tgcagagacc	aacacaatgg	3300
ttcccaagac	aactcccaag	ttttcccata	gtaagtcaga	caccacactc	ccagtagcca	3360
tcaccagtcc	tgggccagaa	gccagttcag	ctgtttcaac	gacaactatc	tcacctgata	3420
tgtcagatct	ggtgacctca	ctggtcccta	gttctgggac	agacaccagt	acaaccttcc	3480
caacattgag	tgagacccca	tatgaaccag	agactacagt	cacgtggctc	actcatcctg	3540
cagaaaccag	cacaacggtt	tctgggacaa	ttcccaactt	ttcccatagg	ggatcagaca	3600
ctgcaccctc	aatggtcacc	agtcctggag	tagacacgag	gtcaggtgtt	ccaactacaa	3660
ccatcccacc	cagtatacca	ggggtagtga	cctcacaggt	cactagttct	gcaacagaca	3720
ctagtacagc	tattccaact	ttgactcctt	ctcctggtga	accagagacc	acagcctcat	3780
cagctaccca	tcctgggaca	cagactggct	tcactgttcc	aattcggact	gttccctcta	3840
gtgagccaga	tacaatggct	tcctgggtca	ctcatcctcc	acagaccagc	acacctgttt	3900
ccagaacaac	ctccagtttt	tcccatagta	gtccagatgc	cacacctgta	atggccacca	3960
gtcctaggac	agaagccagt	tcagctgtac	tgacaacaat	ctcacctggt	gcaccagaga	4020
tggtgacttc	acagatcact	agttctgggg	cagcaaccag	tacaactgtt	ccaactttga	4080
ctcattctcc	tggtatgcca	gagaccacag	ccttattgag	cacccatccc	agaacaggga	4140
caagtaaaac	atttcctgct	tcaactgtgt	ttcctcaagt	atcagagacc	acagcctcac	4200
tcaccattag	acctggtgca	gagactagca	cagctctccc	aactcagaca	acatcctctc	4260
tcttcaccct	acttgtaact	ggaaccagca	gagttgatct	aagtccaact	gcttcacctg	4320
gtgtttctgc	aaaaacagcc	ccactttcca	cccatccagg	gacagagacc	agcacaatga	4380
ttccaacttc	aactctttcc	cttggtttac	tagagactac	aggcttactg	gccaccagct	4440
cttcagcaga	gaccagcacg	agtactctaa	ctctgactgt	ttcccctgct	gtctctgggc	4500

tttccagtgc	ctctataaca	actgataagc	cccaaactgt	gacctcctgg	aacacagaaa	4560
cctcaccatc	tgtaacttca	gttggacccc	cagaattttc	caggactgtc	acaggcacca	4620
ctatgacctt	gataccatca	gagatgccaa	caccacctaa	aaccagtcat	ggagaaggag	4680
tgagtccaac	cactatcttg	agaactacaa	tggttgaagc	cactaattta	gctaccacag	4740
gttccagtcc	cactgtggcc	aagacaacaa	ccaccttcaa	tacactggct	ggaagcctct	4800
ttactcctct	gaccacacct	gggatgtcca	ccttggcctc	tgagagtgtg	acctcaagaa	4860
caagttataa	ccatcggtcc	tggatctcca	ccaccagcag	ttataaccgt	cggtactgga	4920
cccctgccac	cagcactcca	gtgacttcta	cattctcccc	agggatttcc	acatcctcca	4980
tccccagctc	cacagcagcc	acagtcccat	tcatggtgcc	attcaccctc	aacttcacca	5040
tcaccaacct	gcagtacgag	gaggacatgc	ggcaccctgg	ttccaggaag	ttcaacgcca	5100
cagagagaga	actgcagggt	ctgctcaaac	ccttgttcag	gaatagcagt	ctggaatacc	5160
tctattcagg	ctgcagacta	gcctcactca	ggccagagaa	ggatagctca	gccatggcag	5220
tggatgccat	ctgcacacat	cgccctgacc	ctgaagacct	cggactggac	agagagcgac	5280
tgtactggga	gctgagcaat	ctgacaaatg	gcatccagga	gctgggcccc	tacaccctgg	5340
accggaacag	tctctatgtc	aatggtttca	cccatcgaag	ctctatgccc	accaccagca	5400
ctcctgggac	ctccacagtg	gatgtgggaa	cctcagggac	tccatcctcc	agccccagcc	5460
ccacg						5465

<210> 82 <211> 1821

<212> PRT

<213> Homo sapiens

<400> 82

Glu Ser Val Leu Glu Gly Thr Val Thr Ser Ala Tyr Gln Val Pro Ser 10 15

Leu Ser Thr Arg Leu Thr Arg Thr Asp Gly Ile Met Glu His Ile Thr $20 \hspace{1cm} 25 \hspace{1cm} 30$

Lys Ile Pro Asn Glu Ala Ala His Arg Gly Thr Ile Arg Pro Val Lys 35 40 45

Gly Pro Gln Thr Ser Thr Ser Pro Ala Ser Pro Lys Gly Leu His Thr 50 60

Gly Gly Thr Lys Arg Met Glu Thr Thr Thr Thr Ala Leu Lys Thr Thr 65 70 75 80

Thr Thr Ala Leu Lys Thr Thr Ser Arg Ala Thr Leu Thr Thr Ser Val Page 102

Tyr Thr Pro Thr Leu Gly Thr Leu Thr Pro Leu Asn Ala Ser Arg Gln Met Ala Ser Thr Ile Leu Thr Glu Met Met Ile Thr Thr Pro Tyr Val 115 120 125 Phe Pro Asp Val Pro Glu Thr Thr Ser Ser Leu Ala Thr Ser Leu Gly Ala Glu Thr Ser Thr Ala Leu Pro Arg Thr Thr Pro Ser Val Leu Asn Arg Glu Ser Glu Thr Thr Ala Ser Leu Val Ser Arg Ser Gly Ala Glu Arg Ser Pro Val Ile Gln Thr Leu Asp Val Ser Ser Glu Pro Asp 180 185 190 Thr Thr Ala Ser Trp Val Ile His Pro Ala Glu Thr Ile Pro Thr Val 195 200 205 Ser Lys Thr Thr Pro Asn Phe Phe His Ser Glu Leu Asp Thr Val Ser 210 220 Ser Thr Ala Thr Ser His Gly Ala Asp Val Ser Ser Ala Ile Pro Thr 225 230 235 240 Asn Ile Ser Pro Ser Glu Leu Asp Ala Leu Thr Pro Leu Val Thr Ile 245 250 255 Ser Gly Thr Asp Thr Ser Thr Thr Phe Pro Thr Leu Thr Lys Ser Pro 260 265 270 His Glu Thr Glu Thr Arg Thr Thr Trp Leu Thr His Pro Ala Glu Thr Ser Ser Thr Ile Pro Arg Thr Ile Pro Asn Phe Ser His His Glu Ser 290 295 300 Asp Ala Thr Pro Ser Ile Ala Thr Ser Pro Gly Ala Glu Thr Ser Ser 305 310 315 320 Ala Ile Pro Ile Met Thr Val Ser Pro Gly Ala Glu Asp Leu Val Thr 325 330 335

09-965738substitute.ST25.txt Ser Gln Val Thr Ser Ser Gly Thr Asp Arg Asn Met Thr Ile Pro Thr Leu Thr Leu Ser Pro Gly Glu Pro Lys Thr Ile Ala Ser Leu Val Thr His Pro Glu Ala Gln Thr Ser Ser Ala Ile Pro Thr Ser Thr Ile Ser Pro Ala Val Ser Arg Leu Val Thr Ser Met Val Thr Ser Leu Ala Ala Lys Thr Ser Thr Thr Asn Arg Ala Leu Thr Asn Ser Pro Gly Glu Pro Ala Thr Thr Val Ser Leu Val Thr His Pro Ala Gln Thr Ser Pro Thr Val Pro Trp Thr Thr Ser Ile Phe Phe His Ser Lys Ser Asp Thr Thr 440 Pro Ser Met Thr Thr Ser His Gly Ala Glu Ser Ser Ser Ala Val Pro Thr Pro Thr Val Ser Thr Glu Val Pro Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Ile Ser Thr Thr Ile Pro Ile Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Glu Glu Ala Ser Ser Ala Ile Pro Thr Pro Thr Val Ser Pro Gly Val
515 520 525 Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr Phe Ser Leu Gly Glu Pro Glu Thr Thr 545 550 555 560 545 Pro Ser Met Ala Thr Ser His Gly Thr Glu Ala Gly Ser Ala Val Pro Thr Val Leu Pro Glu Val Pro Gly Met Val Thr Ser Leu Val Ala Ser 580

Ser Arg Ala Val Thr Ser Thr Thr Leu Pro Thr Leu Thr Leu Ser Pro 595 Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Thr Val Pro Thr Val Ser Pro Glu Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Ser Gly Val Asn Ser Thr Ser Ile Pro Thr Leu Ile Leu Ser Pro Gly Glu Leu Glu Thr Thr Pro Ser Met Ala 660 Thr Ser His Gly Ala Glu Ala Ser Ser Ala Val Pro Thr Pro Thr Val Ser Pro Gly Val Ser Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr Leu Ser Ser Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Val Glu Ala Ser 725 730 735 Ser Ala Val Leu Thr Val Ser Pro Glu Val Pro Gly Met Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Thr Leu 755 760 765 Thr Ile Ser Ser Asp Glu Pro Glu Thr Thr Thr Ser Leu Val Thr His Ser Glu Ala Lys Met Ile Ser Ala Ile Pro Thr Leu Ala Val Ser Pro Thr Val Gln Gly Leu Val Thr Ser Leu Val Thr Ser Ser Gly Ser Glu Thr Ser Ala Phe Ser Asn Leu Thr Val Ala Ser Ser Gln Pro Glu Thr Ile Asp Ser Trp Val Ala His Pro Gly Thr Glu Ala Ser Ser Val Val 840

Page 105

- Pro Thr Leu Thr Val Ser Thr Gly Glu Pro Phe Thr Asn Ile Ser Leu 850 855 860
- Val Thr His Pro Ala Glu Ser Ser Ser Thr Leu Pro Arg Thr Thr Ser 865 870 875 880
- Arg Phe Ser His Ser Glu Leu Asp Thr Met Pro Ser Thr Val Thr Ser 885 890 895
- Pro Glu Ala Glu Ser Ser Ser Ala Ile Ser Thr Thr Ile Ser Pro Gly 900 905 910
- Ile Pro Gly Val Leu Thr Ser Leu Val Thr Ser Ser Gly Arg Asp Ile 915 920 925
- Ser Ala Thr Phe Pro Thr Val Pro Glu Ser Pro His Glu Ser Glu Ala 930 935 940
- Thr Ala Ser Trp Val Thr His Pro Ala Val Thr Ser Thr Thr Val Pro 945 950 955 960
- Arg Thr Thr Pro Asn Tyr Ser His Ser Glu Pro Asp Thr Thr Pro Ser 965 970 975
- Ile Ala Thr Ser Pro Gly Ala Glu Ala Thr Ser Asp Phe Pro Thr Ile 980 985 990
- Thr Val Ser Pro Asp Val Pro Asp Met Val Thr Ser Gln Val Thr Ser 995 1000 1005
- Ser Gly Thr Asp Thr Ser Ile Thr Ile Pro Thr Leu Thr Leu Ser 1010 1015 1020
- Ser Gly Glu Pro Glu Thr Thr Thr Ser Phe Ile Thr Tyr Ser Glu 1025 1030 1035
- Thr His Thr Ser Ser Ala Ile Pro Thr Leu Pro Val Ser Pro Gly 1040 1045 1050
- Ala Ser Lys Met Leu Thr Ser Leu Val Ile Ser Ser Gly Thr Asp 1055 1060 1065
- Ser Thr Thr Thr Phe Pro Thr Leu Thr Glu Thr Pro Tyr Glu Pro 1070 1075 1080
- Glu Thr Thr Ala Ile Gln Leu Ile His Pro Ala Glu Thr Asn Thr Page 106

1085

Met	val 1100	Pro	Arg	Thr	Thr	Pro 1105	Lys	Phe	Ser	His	Ser 1110	Lys	Ser	Asp
Thr	Thr 1115	Leu	Pro	Val	Ala	Ile 1120	Thr	Ser	Pro	Gly	Pro 1125	Glu	Ala	Ser
Ser	Ala 1130	val	Ser	Thr	Thr	Thr 1135	Ile	Ser	Pro	Asp	Met 1140	Ser	Asp	Leu
val	Thr 1145	Ser	Leu	val	Pro	Ser 1150	Ser	Gly	Thr	Asp	Thr 1155	Ser	Thr	Thr
Phe	Pro 1160	Thr	Leu	Ser	Glu	Thr 1165	Pro	Tyr	Glu	Pro	Glu 1170	Thr	Thr	Ala
Thr	Trp 1175	Leu	Thr	His	Pro	Ala 1180	Glu	Thr	Ser	Thr	Thr 1185	val	Ser	Gly
Thr	Ile 1190	Pro	Asn	Phe	Ser	ніs 1195	Arg	Gly	Ser	Asp	Thr 1200	Ala	Pro	Ser
Met	Val 1205	Thr	Ser	Pro	Gly	Val 1210	Asp	Thr	Arg	Ser	Gly 1215	val	Pro	Thr
Thr	Thr 1220	Ile	Pro	Pro	Ser	Ile 1225	Pro	Gly	٧a٦	val	Thr 1230	Ser	Gln	val
Thr	Ser 1235	Ser	Ala	Thr	Asp	Thr 1240	Ser	Thr	Ala	Ile	Pro 1245	Thr	Leu	Thr
Pro	Ser 1250	Pro	Gly	Glu	Pro	Glu 1255	Thr	Thr	Ala	Ser	Ser 1260	Ala	Thr	His
Pro	Gly 1265	Thr	Gln	Thr	Gly	Phe 1270	Thr	val	Pro	Ile	Arg 1275	Thr	val	Pro
Ser	Ser 1280	Glu	Pro	Asp	Thr	Met 1285	Ala	Ser	Trp	val	Thr 1290	ніѕ	Pro	Pro
Gln	Thr 1295	Ser	Thr	Pro	val	Ser 1300	Arg	Thr	Thr	Ser	Ser 1305	Phe	Ser	His
Ser	Ser 1310	Pro	Asp	Ala	Thr	Pro 1315	۷al	Met	Ala	Thr	Ser 1320	Pro	Arg	Thr

Glu	Ala 1325	Ser	Ser	Ala	val	09 Leu 1330	Thr				re.ST2 Pro 1335			Pro
Glu	Met 1340	val	Thr	Ser	Gln	Ile 1345	Thr	Ser	Ser	Gly	Ala 1350	Ala	Thr	Ser
Thr	Thr 1355	val	Pro	Thr	Leu	Thr 1360	His	Ser	Pro	Gly	Met 1365	Pro	Glu	Thr
Thr	Ala 1370	Leu	Leu	Ser	Thr	Нis 1375	Pro	Arg	Thr	Glu	Thr 1380	Ser	Lys	Thr
Phe	Pro 1385	Ala	Ser	Thr	٧a٦	Phe 1390	Pro	Gln	val	Ser	Glu 1395	Thr	Thr	Ala
Ser	Leu 1400	Thr	Ile	Arg	Pro	Gly 1405	Ala	Glu	Thr	Ser	Thr 1410	Ala	Leu	Pro
Thr	Gln 1415	Thr	Thr	Ser	Ser	Leu 1420	Phe	Thr	Leu	Leu	Val 1425	Thr	Gly	Thr
Ser	Arg 1430	۷al	Asp	Leu	Ser	Pro 1435	Thr	Ala	Ser	Pro	Gly 1440	val	Ser	Ala
Lys	Thr 1445	Ala	Pro	Leu	Ser	Thr 1450	His	Pro	Gly	Thr	Glu 1455	Thr	Ser	Thr
Met	Ile 1460	Pro	Thr	Ser	Thr	Leu 1465	Ser	Leu	Gly	Leu	Leu 1470	Glu	Thr	Thr
Gly	Leu 1475	Leu	Аlа	Thr	Ser	Ser 1480	Ser	Ala	Glu	Thr	Ser 1485	Thr	Ser	Thr
Leu	Thr 1490	Leu	Thr	Val	Ser	Pro 1495	Ala	٧a٦	Ser	Gly	Leu 1500	Ser	Ser	Ala
Ser	Ile 1505	Thr	Thr	Asp	Lys	Pro 1510	Gln	Thr	٧a٦	Thr	Ser 1515	Trp	Asn	Thr
Glu	Thr 1520	Ser	Pro	Ser	val	Thr 1525	Ser	val	Gly	Pro	Pro 1530	Glu	Phe	Ser
Arg	Thr 1535	val	Thr	Gly	Thr	Thr 1540	Met	Thr	Leu	Ile	Pro 1545	Ser	Glu	Met
Pro	Thr 1550	Pro	Pro	Lys	Thr	Ser 1555	Нis	Gly			1560	Ser	Pro	Thr
									Page	e 10	8			

Thr Ile Leu Arg Thr Thr Met Val Glu Ala Thr Asn_ Leu Ala Thr 1565 Thr Gly Ser Ser Pro Thr Val Ala Lys Thr Thr Thr Phe Asn 1580 1585 1590 Thr Leu Ala Gly Ser Leu Phe Thr Pro Leu Thr Thr Pro Gly Met 1600 Ser Thr Leu Ala Ser Glu Ser Val Thr Ser Arg Thr Ser Tyr Asn 1620 1615 His Arg Ser Trp Ile Ser Thr Thr Ser Ser Tyr Asn Arg Arg Tyr 1630 Trp Thr Pro Ala Thr Ser Thr Pro Val Thr Ser Thr Phe Ser Pro Gly Ile Ser Thr Ser Ser Ile Pro Ser Ser Thr Ala Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn **16**95 **1690** 1685 Ala Thr Glu Arg Glu Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg 1700 Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser 1715 1720 1725 Leu Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu 1745 1750 1755 Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu 1760 1770 Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 1775 1780 1785 Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr Pro Gly Thr 1795 Page 109

Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser Ser Pro 1805 1810 1815	
Ser Pro Thr 1820	
<210> 83 <211> 468 <212> DNA <213> Homo sapiens	
<400> 83 gccacagtcc cattcatggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac	60
gaggaggaca tgcggcaccc tggttccagg aagttcaacg ccacagagag agaactgcag	120
ggtctgctca aacccttgtt caggaatagc agtctggaat acctctattc aggctgcaga	180
ctagcctcac tcaggccaga gaaggatagc tcagccatgg cagtggatgc catctgcata	240
catcgccctg accctgaaga cctcggactg gacagagagc gactgtactg ggagctgagc	300
aatctgacaa atggcatcca ggagctgggc ccctacaccc tggaccggaa cagtctctat	360
gtcaatggtt tcacccatcg aagctctatg cccaccacca gcactcctgg gacctccaca	420
gtggatgtgg gaacctcagg gactccatcc tccagcccca gccccacg	468
<210> 84 <211> 474 <212> DNA <213> Homo sapiens	
<400> 84 gctgctggcc ctctcctgat gccgttcacc ctcaacttca ccatcaccaa cctgcagtac	60
gaggaggaca tgcgtcgcac tggctccagg aagttcaaca ccatggagag tgtcctgcag	120
ggtctgctca agcccttgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ttgaccttgc tcaggcccaa gaaagatggg gcagccactg gagtggatgc catctgcacc	240
caccgccttg accccaaaag ccctggactc aacagggagc agctgtactg ggagctaagc	300
aaactgacca atgacattga agagctgggc ccctacaccc tggacaggaa cagtctctat	360
gtcaatggtt tcacccatca gagctctgtg tccaccacca gcactcctgg gacctccaca	420
gtggatctca gaacctcagg gactccatcc tccctctcca gccccacaat tatg	474
<210> 85 <211> 468 <212> DNA <213> Homo sapiens	
<400> 85 gctgctggcc ctctcctggt accattcacc ctcaacttca ccatcaccaa cctgcagtat Page 110	60

ggggaggaca tgggtcaccc	tggctccagg	aagttcaaca	ccacagagag	ggtcctgcag	120
ggtctgcttg gtcccatatt	caagaacacc	agtgttggcc	ctctgtactc	tggctgcaga	180
ctgacctctc tcaggtctga	gaaggatgga	gcagccactg	gagtggatgc	catctgcatc	240
catcatcttg accccaaaag	ccctggactc	aacagagagc	ggctgtactg	ggagctgagc	300
caactgacca atggcatcaa	agagctgggc	ccctacaccc	tggacaggaa	cagtctctat	360
gtcaatggtt tcacccatcg	gacctctgtg	cccaccacca	gcactcctgg	gacctccaca	420
gtggaccttg gaacctcagg	gactccattc	tccctcccaa	gccccgca		468
<210> 86 <211> 465 <212> DNA <213> Homo sapiens					
<400> 86 actgctggcc ctctcctggt	gctgttcacc	ctcaacttca	ccatcaccaa	cctgaagtat	60
gaggaggaca tgcatcgccc					120
actctgcttg gtcctatgtt	caagaacacc	agtgttggcc	ttctgtactc	tggctgcaga	180
ctgaccttgc tcaggtccga	gaaggatgga	gcagccactg	gagtggatgc	catctgcacc	240
caccgtcttg accccaaaag	ccctggactg	gacagagagc	agctatactg	ggagctgagc	300
cagctgacca atggcatcaa	agagctgggc	ccctacaccc	tggacaggaa	cagtctctat	360
gtcaatggtt tcacccattg	gatccctgtg	cccaccagca	gcactcctgg	gacctccaca	420
gtggaccttg ggtcagggac	tccatcctcc	ctccccagcc	ccaca		465
<210> 87 <211> 468 <212> DNA <213> Homo sapiens					
<400> 87 gctgctggcc ctctcctggt	gccattcacc	ctcaacttca	ccatcaccaa	cctgcagtac	60
gaggaggaca tgcatcaccc	aggctccagg	aagttcaaca	ccacggagcg	ggtcctgcag	120
ggtctgcttg gtcccatgtt	caagaacacc	agtgtcggcc	ttctgtactc	tggctgcaga	180
ctgaccttgc tcaggtccga	gaaggatgga	gcagccactg	gagtggatgc	catctgcacc	240
caccgtcttg accccaaaag	ccctggagtg	gacagggagc	agctatactg	ggagctgagc	300
cagctgacca atggcatcaa	agagctgggt	ccctacaccc	tggacagaaa	cagtctctat	360
gtcaatggtt tcacccatca	gacctctgcg	cccaacacca	gcactcctgg	gacctccaca	420
gtggaccttg ggacctcagg	gactccatcc	tccctcccca	gccctaca		468

09-965738substitute.ST25.txt	
<211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 88 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan	60
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag	120
ggtctgctnn nncccntntt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ctgaccttgc tcaggtccga gaaggatgga gcagccactg gagtggatgc catctgcacc	240
caccgtcttg accccaaaag ccctggagtg gacagggagc agctatactg ggagctgagc	300
cagctgacca atggcatcaa agagctgggt ccctacaccc tggacagaaa cagtctctat	360
gtcaatggtt tcacccatca gacctctgcg cccaacacca gcactcctgg gacctccaca	420
gtggaccttg ggacctcagg gactccatcc tccctcccca gccctaca	468
<210> 89 <211> 468 <212> DNA <213> Homo sapiens	
<400> 89 tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac	60
gaggaggaca tgcatcaccc aggctccagg aagttcaaca ccacggagcg ggtcctgcag	120
ggtctgcttg gtcccatgtt caagaacacc agtgtcggcc ttctgtactc tggctgcaga	180
ctgaccttgc tcaggcctga gaagaatggg gcagccactg gaatggatgc catctgcagc	240
caccgtcttg accccaaaag ccctggactc aacagagagc agctgtactg ggagctgagc	300
cagctgaccc atggcatcaa agagctgggc ccctacaccc tggacaggaa cagtctctat	360
gtcaatggtt tcacccatcg gagctctgtg gcccccacca gcactcctgg gacctccaca	420
gtggaccttg ggacctcagg gactccatcc tccctccca gccccaca	468
<210> 90 <211> 468 <212> DNA <213> Homo sapiens	
<400> 90 acagctgttc ctctcctggt gccgttcacc ctcaacttta ccatcaccaa tctgcagtat	60
ggggaggaca tgcgtcaccc tggctccagg aagttcaaca ccacagagag ggtcctgcag	120
ggtctgcttg gtcccttgtt caagaactcc agtgtcggcc ctctgtactc tggctgcaga	180

ctgatctctc tcaggtctga			tute.ST25.tx gagtggatgc		240
caccacctta accctcaaag					300
cagatgacca atggcatcaa					360
gtcaatggtt tcacccatcg					420
gttgaccttg gaacctcagg	gactccatcc	cccgtcccca	gccccaca		468
		-			
<210> 91 <211> 468 <212> DNA <213> Homo sapiens					
<400> 91					
actgctggcc ctctcctggt	gccattcacc	ctcaacttca	ccatcaccaa	cctgcagtat	60
gaggaggaca tgcatcgccc	tggatctagg	aagttcaaca	ccacagagag	ggtcctgcag	120
ggtctgctta gtcccatttt	caagaactcc	agtgttggcc	ctctgtactc	tggctgcaga	180
ctgacctctc tcaggcccga	gaaggatggg	gcagcaactg	gaatggatgc	tgtctgcctc	240
taccacccta atcccaaaag	acctggactg	gacagagagc	agctgtactg	ggagctaagc	300
cagctgaccc acaacatcac	tgagctgggc	ccctacagcc	tggacaggga	cagtctctat	360
gtcaatggtt tcacccatca	gaactctgtg	cccaccacca	gtactcctgg	gacctccaca	420
gtgtactggg caaccactgg	gactccatcc	tccttccccg	gccacaca		468
<210> 92 <211> 468 <212> DNA <213> Homo sapiens					
<400> 92 gagcctggcc ctctcctgat	accattcact	ttcaacttta	ccatcaccaa	cctgcattat	60
gaggaaaaca tgcaacaccc	tggttccagg	aagttcaaca	ccacggagag	ggttctgcag	120
ggtctgctca agcccttgtt	caagaacacc	agtgttggcc	ctctgtactc	tggctgcaga	180
ctgacctctc tcaggcccga	gaaggatggg	gcagcaactg	gaatggatgc	tgtctgcctc	240
taccacccta atcccaaaag	acctgggctg	gacagagagc	agctgtactg	ggagctaagc	300
cagctgaccc acaacatcac	tgagctgggc	ccctacagcc	tggacaggga	cagtctctat	360
gtcaatggtt tcacccatca	gaactctgtg	cccaccacca	gtactcctgg	gacctccaca	420
gtgtactggg caaccactgg	gactccatcc	tccttccccg	gccacaca		468
<210> 93 <211> 468 <212> DNA <213> Homo sapiens					

<400> 93

00.055700 1.1.1.1.1.1.1.1.1.1.1.1.1.1.1.1.1.1.1	
09-965738substitute.ST25.txt gagcctggcc ctctcctgat accattcact ttcaacttta ccatcaccaa cctgcattat	60
gaggaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag	120
ggtctgctca agcccttgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ctgaccttgc tcagacctga gaagcatgag gcagccactg gagtggacac catctgtacc	240
caccgcgttg atcccatcgg acctggactg gacagggagc ggctatactg ggagctgagc	300
cagctgacca acagcattac cgaactggga ccctacaccc tggacaggga cagtctctat	360
gtcaatggct tcaaccctcg gagctctgtg ccaaccacca gcactcctgg gacctccaca	420
gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca	468
<210> 94 <211> 468 <212> DNA <213> Homo sapiens	
<pre><220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide</pre>	
<400> 94 gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat	60
gaggaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag	120
ggtctgctca agcccttgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ctgaccttgc tcagacctga gaagcatgag gcagccactg gagtggacac catctgtacc	240
caccgcgttg atcccatcgg acctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca	420
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca	468
<210> 95 · · · · · · · · · · · · · · · · · ·	
<400> 95 tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac	60
gaggaggaca tgcatcaccc aggctccagg aagttcaaca ccacggagcg ggtcctgcag	120
ggtctgcttg gtcccatgtt caagaacacc agtgtcggcc ttctgtactc tggctgcaga	180
ctgaccttgc tcaggcctga gaagaatggg gcagccactg gaatggatgc catctgcagc	240
caccgtcttg accccaaaag ccctggactc gacagagagc agctgtactg ggagctgagc	300
cagctgaccc atggcatcaa agagctgggc ccctacaccc tggacaggaa cagtctctat Page 114	360

gtcaatggtt tcacccatcg gagctctgtg gcccccacca gcactcctgg gacctccaca	420
gtggaccttg ggacctcagg gactccatcc tccctcccca gccccaca	468
<210> 96 <211> 468 <212> DNA <213> Homo sapiens	
<400> 96 acagctgttc ctctcctggt gccgttcacc ctcaacttta ccatcaccaa tctgcagtat	60
ggggaggaca tgcgtcaccc tggctccagg aagttcaaca ccacagagag ggtcctgcag	120
ggtctgcttg gtcccttgtt caagaactcc agtgtcggcc ctctgtactc tggctgcaga	180
ctgatctctc tcaggtctga gaaggatggg gcagccactg gagtggatgc catctgcacc	240
caccacctta accctcaaag ccctggactg gacagggagc agctgtactg gcagctgagc	300
cagatgacca atggcatcaa agagctgggc ccctacaccc tggaccggaa cagtctctac	360
gtcaatggtt tcacccatcg gagctctggg ctcaccacca gcactccttg gacttccaca	420
gttgaccttg gaacctcagg gactccatcc cccgtcccca gccccaca	468
<210> 97 <211> 468 <212> DNA <213> Homo sapiens	
<400> 97 actgctggcc ctctcctggt gccattcacc ctaaacttca ccatcaccaa cctgcagtat	60
gaggaggaca tgcatcgccc tggatctagg aagttcaacg ccacagagag ggtcctgcag	120
gaggaggaca tgcatcgccc tggatctagg aagttcaacg ccacagagag ggtcctgcag ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga	120 180
ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga	180
ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga ctgacctctc tcaggcccga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc	180 240
ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga ctgacctctc tcaggcccga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc taccacccta atcccaaaag acctggactg gacagagagc agctgtactg ggagctaagc	180 240 300
ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga ctgacctctc tcaggcccga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc taccacccta atcccaaaag acctggactg gacagagagc agctgtactg ggagctaagc cagctgaccc acaacatcac tgagctgggc ccctacagcc tggacaggga cagtctctat	180 240 300 360
ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga ctgacctctc tcaggcccga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc taccacccta atcccaaaag acctggactg gacagagagc agctgtactg ggagctaagc cagctgaccc acaacatcac tgagctgggc ccctacagcc tggacaggga cagtctctat gtcaatggtt tcacccatca gagctctatg acgaccacca gaactcctga tacctccaca	180 240 300 360 420
ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga ctgacctctc tcaggcccga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc taccacccta atcccaaaag acctggactg gacagagagc agctgtactg ggagctaagc cagctgaccc acaacatcac tgagctgggc ccctacagcc tggacaggga cagtctctat gtcaatggtt tcacccatca gagctctatg acgaccacca gaactcctga tacctccaca atgcacctgg caacctcgag aactccagcc tccctgtctg gacctacg	180 240 300 360 420
ggtctgctta gtcccatatt caagaactcc agtgttggcc ctctgtactc tggctgcaga ctgacctctc tcaggcccga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc taccacccta atcccaaaag acctggactg gacagagagc agctgtactg ggagctaagc cagctgaccc acaacatcac tgagctgggc ccctacagcc tggacaggga cagtctctat gtcaatggtt tcacccatca gagctctatg acgaccacca gaactcctga tacctccaca atgcacctgg caacctcgag aactccagcc tccctgtctg gacctacg <210> 98 <211> 474 <212> DNA <213> Homo sapiens <400> 98	180 240 300 360 420 468

ttgaccttgc tcaggcccaa gaaagatggg gcagccactg gagtggatgc catctgcacc	240
caccgccttg accccaaaag ccctggactc aacagggagc agctgtactg ggagctaagc	300
aaactgacca atgacattga agagctgggc ccctacaccc tggacaggaa cagtctctat	360
gtcaatggtt tcacccatca gagctctgtg tccaccacca gcactcctgg gacctccaca	420
gtggatctca gaacctcagg gactccatcc tccctctcca gccccacaat tatg	474
graduction gamestoday garetodates totales a graduation care	
<210> 99 <211> 468	
<212> DNA <213> Homo sapiens	
TELEST TIOMS SUPPLIES	
<220> <221> misc_feature	
<222> (1)(468) <223> All N's = any nucleotide	
<400> 99	
ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan	60
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagag ggtcctacag	120
ggtctgctca ggcccttgtt caagaacacc agtgtcagct ctctgtactc tggttgcaga	180
ctgaccttgc tcaggcctga gaaggatggg gcagccacca gagtggatgc tgcctgcacc	240
taccgccctg atcccaaaag ccctggactg gacagagagc aactatactg ggagctgagc	300
cagctaaccc acagcatcac tgagctggga ccctacaccc tggacagggt cagtctctat	360
gtcaatggct tcaaccctcg gagctctgtg ccaaccacca gcactcctgg gacctccaca	420
gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca	468
<210> 100	
<211> 468 <212> DNA	
<213> Homo sapiens	
<400> 100 gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat	60
gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag	120
	180
ggtctgctca agcccttgtt caagagcacc agcgttggcc ctctgtactc tggctgcaga	240
ctgaccttgc tcagacctga gaaacatggg gcagccactg gagtggacgc catctgcacc	300
ctccgccttg atcccactgg tcctggactg gacagagagc ggctatactg ggagctgagc	360
cagctgacca acagcgttac agagctgggc ccctacaccc tggacaggga cagtctctat	420
gtcaatggct tcacccagcg gagctctgtg ccaaccacca gtattcctgg gacctctgca	
gtgcacctgg aaacctctgg gactccagcc tccctccctg gccacaca	468

03 303/303450-144-07-07-07-07-07-07-07-07-07-07-07-07-07-	
<210> 101 <211> 468 <212> DNA <213> Homo sapiens	
<400> 101 gcccctggcc ctctcctggt gccattcacc ctcaacttca ctatcaccaa cctgcagtat	60
gaggtggaca tgcgtcaccc tggttccagg aagttcaaca ccacggagag agtcctgcag 12	20
ggtctgctca agcccttgtt caagagcacc agtgttggcc ctctgtactc tggctgcaga 18	80
ctgaccttgc tcaggcctga aaaacgtggg gcagccaccg gcgtggacac catctgcact 24	40
caccgccttg accctctaaa ccctggactg gacagagagc agctatactg ggagctgagc 30	00
aaactgaccc gtggcatcat cgagctgggc ccctacctcc tggacagagg cagtctctat 36	60
gtcaatggtt tcacccatcg gaactttgtg cccatcacca gcactcctgg gacctccaca 42	20
gtacacctag gaacctctga aactccatcc tccctaccta gacccata 46	68
<210> 102 <211> 468 <212> DNA <213> Homo sapiens	
<400> 102 gtgcctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cttgcagtat	60
gaggaggcca tgcgacaccc tggctccagg aagttcaata ccacggagag ggtcctacag 12	20
ggtctgctca ggcccttgtt caagaatacc agtatcggcc ctctgtactc cagctgcaga 18	80
ctgaccttgc tcaggccaga gaaggacaag gcagccacca gagtggatgc catctgtacc 24	40
caccacctg accctcaaag ccctggactg aacagagagc agctgtactg ggagctgagc 30	00
cagctgaccc acggcatcac tgagctgggc ccctacaccc tggacaggga cagtctctat 36	60
gtcgatggtt tcactcattg gagccccata ccgaccacca gcactcctgg gacctccata 42	20
gtgaacctgg gaacctctgg gatcccacct tccctcctg aaactaca 46	68
<210> 103 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 103 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan	60
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagag ggttctgcag 12	20

			ute.ST25.tx		400
ggtctgctca aacccttgtt					180
ctagcctcac tcaggccaga	gaaggatagc	tcagccatgg	cagtggatgc	catctgcaca	240
catcgccctg accctgaaga	cctcggactg	gacagagagc	gactgtactg	ggagctgagc	300
aatctgacaa atggcatcca	ggagctgggc	ccctacaccc	tggaccggaa	cagtctctac	360
gtcaatggtt tcacccatcg	gagctctggg	ctcaccacca	gcactccttg	gacttccaca	420
gttgaccttg gaacctcagg	gactccatcc	cccgtcccca	gccccaca		468
<210> 104 <211> 468 <212> DNA <213> Homo sapiens					
<400> 104 actgctggcc ctctcctggt	gccattcacc	ctcaacttca	ccatcaccaa	cctgcagtat	60
gaggaggaca tgcatcgccc	tggttccagg	aggttcaaca	ccacggagag	ggttctgcag	120
ggtctgctca cgcccttgtt	caagaacacc	agtgttggcc	ctctgtactc	tggctgcaga	180
ctgaccttgc tcagacctga	gaagcaagag	gcagccactg	gagtggacac	catctgtacc	240
caccgcgttg atcccatcgg	acctggactg	gacagagagc	ggctatactg	ggagctgagc	300
cagctgacca acagcatcac	agagctggga	ccctacaccc	tggataggga	cagtctctat	360
gtcaatggct tcaacccttg	gagctctgtg	ccaaccacca	gcactcctgg	gacctccaca	420
gtgcacctgg caacctctgg	gactccatcc	tccctgcctg	gccacaca		468
<210> 105 <211> 468 <212> DNA <213> Homo sapiens					
<400> 105 gcccctgtcc ctctcttgat	accattcacc	ctcaacttta	ccatcaccga	cctgcattat	60
gaagaaaaca tgcaacaccc	tggttccagg	aagttcaaca	ccacggagag	ggttctgcag	120
ggtctgctca agcccttgtt	caagagcacc	agcgttggcc	ctctgtactc	tggctgcaga	180
ctgaccttgc tcagacctga	gaaacatggg	gcagccactg	gagtggacgc	catctgcacc	240
ctccgccttg atcccactgg	tcctggactg	gacagagagc	ggctatactg	ggagctgagc	300
cagctgacca acagcgttac	agagctgggc	ccctacaccc	tggacaggga	cagtctctat	360
gtcaatggct tcacccatcg	gagctctgtg	ccaaccacca	gtattcctgg	gacctctgca	420
gtgcacctgg aaacctctgg	gactccagcc	tccctccctg	gccacaca		468
<210> 106 <211> 468 <212> DNA <213> Homo sapiens		Page 1	118		

<400> 106					
gcccctggcc ctctcctggt	gccattcacc	ctcaacttca	ctatcaccaa	cctgcagtat	60
gaggaggaca tgcgtcaccc	tggttccagg	aagttcagca	ccacggagag	agtcctgcag	120
ggtctgctca agcccttgtt	caagaacacc	agtgtcagct	ctctgtactc	tggttgcaga	180
ctgaccttgc tcaggcctga	gaaggatggg	gcagccacca	gagtggatgc	tgtctgcacc	240
catcgtcctg accccaaaag	ccctggactg	gacagagagc	ggctgtactg	gaagctgagc	300
cagctgaccc acggcatcac	tgagctgggc	ccctacaccc	tggacaggca	cagtctctat	360
gtcaatggtt tcacccatca	gagctctatg	acgaccacca	gaactcctga	tacctccaca	420
atgcacctgg caacctcgag	aactccagcc	tccctgtctg	gacctacg		468
<210> 107 <211> 468 <212> DNA <213> Homo sapiens <400> 107			,		
accgccagcc ctctcctggt	gctattcaca	attaacttca	ccatcactaa	cctgcggtat	60
gaggagaaca tgcatcaccc	tggctctaga	aagtttaaca	ccacggagag	agtccttcag	120
ggtctgctca ggcctgtgtt	caagaacacc	agtgttggcc	ctctgtactc	tggctgcaga	180
ctgaccacgc tcaggcccaa	gaaggatggg	gcagccacca	aagtggatgc	catctgcacc	240
taccgccctg atcccaaaag	ccctggactg	gacagagagc	agctatactg	ggagctgagc	300
cagctaaccc acagcatcac	tgagctgggc	ccctacaccc	aggacaggga	cagtctctat	360
gtcaatggct tcacccatcg	gagctctgtg	ccaaccacca	gtattcctgg	gacctctgca	420
gtgcacctgg aaacctctgg	gactccagcc	tccctccctg	gccacaca		468
<210> 108 <211> 468 <212> DNA <213> Homo sapiens <400> 108					
gcccctggcc ctctcctggt	gccattcacc	ctcaacttca	ctatcaccaa	cctgcagtat	60
gaggaggaca tgcgtcaccc	tggttccagg	aagttcaaca	ccacggagag	agtcctgcag	120
ggtctgctca agcccttgtt	caagagcacc	agtgttggcc	ctctgtactc	tggctgcaga	180
ctgaccttgc tcaggcctga	aaaacgtggg	gcagccaccg	gcgtggacac	catctgcact	240
caccgccttg accctctaaa	cccaggactg	gacagagagc	agctatactg	ggagctgagc	300
aaactgaccc gtggcatcat	cgagctgggc	ccctacctcc	tggacagagg	cagtctctat	360
gtcaatggtt tcacccatcg	gacctctgtg	cccaccacca	gcactcctgg	gacctccaca	420
gtggaccttg gaacctcagg	gactccattc	tccctcccaa Page	gccccgca 119		468

```
<210>
      109
<211>
      465
<212>
      DNA
<213>
      Homo sapiens
<220>
<221>
      misc_feature
<222> (1)..(465)
<223> All N's = any nucleotide
<400> 109
                                                                      60
ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagag ggtcctgcag
                                                                     120
                                                                     180
actctqcttq qtcctatqtt caaqaacacc agtgttggcc ttctgtactc tggctgcaga
                                                                     240
ctgaccttgc tcaggtccga gaaggatgga gcagccactg gagtggatgc catctgcacc
                                                                     300
caccgtcttg accccaaaag ccctggagtg gacagggagc aactatactg ggagctgagc
cagctgacca atggcattaa agaactgggc ccctacaccc tggacaggaa cagtctctat
                                                                     360
gtcaatgggt tcacccattg gatccctgtg cccaccagca gcactcctgg gacctccaca
                                                                     420
                                                                     465
gtggaccttg ggtcagggac tccatcctcc ctccccagcc ccaca
<210>
      110
<211>
       468
<212>
      DNA
<213>
      Homo sapiens
<400> 110
actgctggcc ctctcctggt gccgttcacc ctcaacttca ccatcaccaa cctgaagtac
                                                                       60
                                                                     120
gaggaggaca tgcattgccc tggctccagg aagttcaaca ccacagagag agtcctgcag
                                                                     180
agtctgcttg gtcccatgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga
ctgaccttgc tcaggtccga gaaggatgga gcagccactg gagtggatgc catctgcacc
                                                                     240
                                                                      300
caccgtcttg accccaaaag ccctggagtg gacagggagc agctatactg ggagctgagc
cagctgacca atggcatcaa agagctgggt ccctacaccc tggacagaaa cagtctctat
                                                                      360
gtcaatggtt tcacccatca gacctctgcg cccaacacca gcactcctgg gacctccaca
                                                                      420
                                                                      468
gtggaccttg ggacctcagg gactccatcc tccctcccca gccctaca
<210>
       111
<211>
       465
<212>
      DNA
<213> Homo sapiens
<220>
       misc_feature
<221>
<222> (1)..(465)
```

09-965738substitute.ST25.txt <223> All N's = any nucleotide <400> 111 nonnctance etetectant neentteace nteaacttna ccateaceaa cetgeantan 60 qnqqannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag 120 ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga 180 240 ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc caccnncntn ancccaaaaq ncctqqactq nacaqnqaqc ngctntactq ggaqctnagc 300 canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat 360 420 qtcaatqqtt tcacccattq qatccctqtq cccaccagca gcactcctgg gacctccaca 465 gtggaccttg ggtcagggac tccatcctcc ctccccagcc ccaca <210> 112 468 <211> <212> DNA <213> Homo sapiens <220> <221> misc_feature <222> (1)..(468) All N's = any nucleotide <223> <400> 112 actgctggcc ctctcctggt gccgttcacc ctcaacttca ccatcaccaa cctgaagtac 60 120 gaggaggaca tgcattgccc tggctccagg aagttcaaca ccacagagag agtcctgcag agtctgcttg gtcccatgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga 180 ctgacctcgc tcaggtccga gaaggatgga gcagccactg gagtggatgc catctgcacc 240 300 caccgtgttg accccaaaag ccctggagtg gacagggagc agctatactg ggagctgagc 360 cagctgacca atggcatcaa agagctgggt ccctacaccc tggacagaaa cagtctctat gtcaatggtt tcacccatca gacctctgcg cccaacacca gcactcctgg gacctccaca 420 468 qtqnacntnq qnacctcnqq qactccatcc tccntccccn gccncaca <210> 113 468 <211> <212> DNA <213> Homo sapiens <220> misc_feature <221> <222> (1)..(468)<223> All N's = any nucleotide <400> tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac 60

09-965738substitute.ST25.txt	
gaggaggaca tgcatcaccc aggctccagg aagttcaaca ccacggagcg ggtcctgcag	120
ggtctgcttg gtcccatgtt caagaacacc agtgtcggcc ttctgtactc tggctgcaga	180
ctgaccttgc tcaggcctga gaagaatggg gcaaccactg gaatggatgc catctgcacc	240
caccgtcttg accccaaaag ccctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca	420
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca	468
<210> 114 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 114 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan	60
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagag ggttctgcag	120
ggtctgctca aacccttgtt caggaatagc agtctggaat acctctattc aggctgcaga	180
ctagcctcac tcaggccaga gaaggatagc tcagccatgg cagtggatgc catctgcaca	240
catcgccctg accctgaaga cctcggactg gacagagagc gactgtactg ggagctgagc	300
aatctgacaa atggcatcca ggagctgggc ccctacaccc tggaccggaa cagtctctat	360
gtcaatggtt tcacccatcg aagctctatg cccaccacca gcactcctgg gacctccaca	420
gtggatgtgg gaacctcagg gactccatcc tccagcccca gccccacg	468
<210> 115 <211> 468 <212> DNA <213> Homo sapiens	
<400> 115 actgctggcc ctctcctgat accattcacc ctcaacttca ccatcaccaa cctgcagtat	60
ggggaggaca tgggtcaccc tggctccagg aagttcaaca ccacagagag ggtcctgcag	120
ggtctgcttg gtcccatatt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ctgacctctc tcaggtctga gaaggatgga gcagccactg gagtggatgc catctgcatc	240
catcatcttg accccaaaag ccctggactc aacagagagc ggctgtactg ggagctgagc	300
caactgacca atggcatcaa agagctgggc ccctacaccc tggacaggaa cagtctctat	360
gtcaatggtt tcacccatcg gacctctgtg cccaccacca gcactcctgg gacctccaca Page 122	420

gtggaccttg	gaacctcagg	gactccattc	tccctcccaa	gccccgca		468
<210> 116 <211> 468 <212> DNA <213> Hom						
<222> (1)	c_feature (468) N's = any 1	nucleotide				
<400> 116	; : ctctcctggt	gctgttcacc	ctcaacttca	ccatcaccaa	cctgaagtat	60
gaggaggaca	tgcatcgccc	tggctccagg	aagttcaaca	ccactgagag	ggtcctgcag	120
actctgcttg	gtcctatgtt	caagaacacc	agtgttggcc	ttctgtactc	tggctgcaga	180
ctgaccttgc	tcaggtccga	gaaggatgga	gcagccactg	gagtggatgc	catctgcacc	240
caccgtcttg	accccaaaag	ccctggactg	nacagngagc	ngctntactg	ggagctnagc	300
canctgacca	annncatcnn	ngagctgggn	ccctacaccc	tggacaggna	cagtctctat	360
gtcaatggtt	tcacccatcn	ganctctgng	cccaccacca	gcactcctgg	gacctccaca	420
gtgnacntng	gnacctcngg	gactccatcc	tccntccccn	gccncaca		468
<210> 117 <211> 468 <212> DNA <213> Hom	}				·	
<222> (1)	sc_feature (468) N's = any (nucleotide				
<400> 117	, ctctcctgnt	nccnttcacc	ntcaacttna	ccatcaccaa	cctocantan	60
	tgcnncnccc					120
	ggcctgtgtt					180
						240
	tcaggcccaa					300
	atcccaaaag acagcatcac					360
						420
	tcacccatcg				gacciciyca	468
grgcaccigg	aaaccactgg	yacıccatcc	tecticety	gccacaca		400

09-965738substitute.ST25.txt	
<211> 468 <212> DNA <213> Homo sapiens	
<400> 118 gagcctggcc ctctcctgat accattcact ttcaacttta ccatcaccaa cctgcgttat	60
gaggaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag	120
ggtctgctca cgcccttgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ctgaccttgc tcagacctga gaagcaggag gcagccactg gagtggacac catctgtacc	240
caccgcgttg atcccatcgg acctggactg gacagagagc ggctatactg ggagctgagc	300
cagctgacca acagcatcac agagctggga ccctacaccc tggataggga cagtctctat	360
gtcgatggct tcaacccttg gagctctgtg ccaaccacca gcactcctgg gacctccaca	420
gtgcacctgg caacctctgg gactccatcc cccctgcctg gccacaca	468
<210> 119 <211> 468 <212> DNA <213> Homo sapiens	
<400> 119 gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccga cctgcattat	60
gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag	120
ggtctgctca agcccttgtt caagagcacc agcgttggcc ctctgtactc tggctgcaga	180
ctgaccttgc tcagacctga gaaacatggg gcagccactg gagtggacgc catctgcacc	240
ctccgccttg atcccactgg tcctggactg gacagagagc ggctatactg ggagctgagc	300
cagctgacca acagcatcac agagctggga ccctacaccc tggataggga cagtctctat	360
gtcaatggct tcaacccttg gagctctgtg ccaaccacca gcactcctgg gacctccaca	420
gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca	468
<210> 120 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 120 actgctggcc ctctcctggt gccgttcacc ctcaacttca ccatcaccaa cctgaagtac	60
gaggaggaca tgcattgccc tggctccagg aagttcaaca ccacagagag agtcctgcag	120
agtctgcatg gtcccatgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180

09-965738substitute.ST25.txt	
ctgaccttgc tcaggtccga gaaggatgga gcagccactg gagtggatgc catctgcacc 240	0
caccgtcttg accccaaaag ccctggactg nacagngagc ngctntactg ggagctnagc 300)
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat 360	0
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca 420	0
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca 468	8
<210> 121 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 121 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan 60	0
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag 120	0
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga 180	0
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc 240	0
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc 300	0
canctgacca acagcatcac agagctggga ccctacaccc tggataggga cagtctctat 360	0
gtcaatggtt tcacccatcg aagctctatg cccaccacca gtattcctgg gacctctgca 420	0
gtgcacctgg aaacctctgg gactccagcc tccctcctg gccacaca 468	8
<210> 122 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 122 gcccctggcc ctctcctggt gccattcacc ctcaacttca ctatcaccaa cctgcagtat 60	0
gaggaggaca tgcgtcaccc tggttccagg aagttcaaca ccacggagag agtcctgcag 120	0
ggtctgctca agcccttgtt caagagcacc agtgttggcc ctctgtactc tggctgcaga 18	0
ctgaccttgc tcaggcctga aaaacgtggg gcagccaccg gcgtggacac catctgcact 24	0
caccgccttg accctctaaa ccctggactg nacagngagc ngctntactg ggagctnagc 30	0
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat 36 Page 125	0

gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca	420
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca	468
<210> 123 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 123 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan	60
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag	120
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga	180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc	240
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360
gtcaatggtt ttcaccctcg gagctctgtg ccaaccacca gcactcctgg gacctccaca	420
gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca	468
<210> 124 <211> 468 <212> DNA <213> Homo sapiens	
<220>	
<pre><221> misc_feature <222> (1)(468) <223> All N's = any nucleotide</pre>	
<222> (1)(468) <223> All N's = any nucleotide <400> 124	60
<222> (1)(468) <223> All N's = any nucleotide <400> 124 gccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat	60 120
<pre><222> (1)(468) <223> All N's = any nucleotide <400> 124 gccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagcg ggtcctgcag</pre>	
<pre><222> (1)(468) <223> All N's = any nucleotide <400> 124 gccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagcg ggtcctgcag ggtctgcttg gtcccatgtt caagaacaca agtgtcggcc ttctgtactc tggctgcaga</pre>	120
<pre><222> (1)(468) <223> All N's = any nucleotide <400> 124 gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagcg ggtcctgcag ggtctgcttg gtcccatgtt caagaacaca agtgtcggcc ttctgtactc tggctgcaga ctgaccttgc tcaggcctga gaagaatggg gcagccactg gaatggatgc catctgcagc</pre>	120 180
<pre><222> (1)(468) <223> All N's = any nucleotide <400> 124 gccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagcg ggtcctgcag ggtctgcttg gtcccatgtt caagaacaca agtgtcggcc ttctgtactc tggctgcaga</pre>	120 180 240
<pre><222> (1)(468) <223> All N's = any nucleotide <400> 124 gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat gaagaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagcg ggtcctgcag ggtctgcttg gtcccatgtt caagaacaca agtgtcggcc ttctgtactc tggctgcaga ctgaccttgc tcaggcctga gaagaatggg gcagccactg gaatggatgc catctgcagc caccgtcttg accccaaaag ccctggactg nacagngagc ngctntactg ggagctnagc</pre>	120 180 240 300

```
<210>
      125
<211>
      468
<212>
      DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222>
      (1)..(468)
All N's = any nucleotide
<223>
<400> 125
ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan
                                                                        60
                                                                       120
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag
                                                                       180
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga
                                                                       240
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc
                                                                       300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat
                                                                       360
gtcaatggtt tcacccatca gaactctgtg cccaccacca gtactcctgg gacctccaca
                                                                       420
gtgtactggg caaccactgg gactccatcc tccttccccg gccacaca
                                                                       468
<210>
      126
<211>
      468
<212>
      DNA
<213> Homo sapiens
<220>
      misc_feature
<221>
<222> (1)..(468)
<223> All N's = any nucleotide
<400>
                                                                        60
gagcctggcc ctctcctgat accattcact ttcaacttta ccatcaccaa cctgcattat
gaggaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag
                                                                       120
                                                                       180
ggtctgctca cgcccttgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga
ctgaccttgc tcagacctga gaagcaggag gcagccactg gagtggacac catctgtacc
                                                                       240
                                                                       300
caccgcqttq atcccatcqq acctggactg nacagngagc ngctntactg ggagctnagc
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat
                                                                       360
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca
                                                                       420
                                                                       468
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca
<210>
       127
<211>
       468
```

:212>

<213>

DNA

Homo sapiens

Page 127

```
<220>
<221> misc_feature
<222> (1)..(468)
<223> All N's = any nucleotide
<400> 127
ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan
                                                                       60
                                                                      120
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag
                                                                      180
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga
                                                                      240
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc
                                                                      300
                                                                      360
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat
gtcaatggtt tcacccatcg gagctctgtg ccaaccacca gcagtcctgg gacctccaca
                                                                      420
gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca
                                                                      468
<210>
       128
<211>
       468
<212>
      DNA
<213>
      Homo sapiens
<220>
<221> misc_feature
<222> (1)..(468)
<223> All N's = any nucleotide
<400> 128
gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcattat
                                                                        60
                                                                       120
qaaqaaaaca tgcaacaccc tggttccagg aagttcaaca ccacggagag ggttctgcag
ggtctgctca agcccttgtt caagagcacc agtgttggcc ctctgtactc tggctgcaga
                                                                       180
ctgaccttgc tcagacctga gaaacatggg gcagccactg gagtggacgc catctgcacc
                                                                       240
ctccgccttg atcccactgg tcctggactg nacagngagc ngctntactg ggagctnagc
                                                                       300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat
                                                                       360
                                                                       420
qtcaatqqtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca
                                                                       468
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca
<210>
       129
<211>
       468
<212>
       DNA
       Homo sapiens
<220>
       misc_feature
<221>
<222> (1)..(468)
```

09-965738substitute.ST25.txt <223> All N's = any nucleotide <400> 129 60 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag 120 ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga 180 ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc 240 300 caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat 360 gtcaatggtt tcacccatcg gacctctgtg cccaccacca gcactcctgg gacctccaca 420 468 gtgcacctgg caacctctgg gactccatcc tccctgcctg gccacaca <210> 130 468 <211> <212> DNA <213> Homo sapiens <220> <221> misc_feature (1)..(468) All N's = any nucleotide <222> <223> <400> gcccctgtcc ctctcttgat accattcacc ctcaacttta ccatcaccaa cctgcagtat 60 120 gaggaggaca tgcatcgccc tggatctagg aagttcaaca ccacagagag ggtcctgcag ggtctgctta gtcccatttt caagaactcc agtgttggcc ctctgtactc tggctgcaga 180 ctgacctctc tcaggcccga gaaggatggg gcagcaactg gaatggatgc tgtctgcctc 240 taccacccta atcccaaaag acctggactg nacagngagc ngctntactg ggagctnagc 300 canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat 360 gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca 420 468 gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca <210> 131 <211> 468 <212> DNA Homo sapiens <213> <220> misc_feature <221> (1)...(468) All N's = any nucleotide <222> <223> <400> ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan 60

Page 129

09-965738substitute.ST25.txt	
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag	120
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga	180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc	240
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360
gtcaatggtt tcacccattg gagctctggg ctcaccacca gcactccttg gacttccaca	420
gttgaccttg gaacctcagg gactccatcc cccgtcccca gccccaca	468
<210> 132 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 132 actgctggcc ctctcctggt gccattcacc ctaaacttca ccatcaccaa cctgcagtat	60
gaggaggaca tgcatcgccc tggatctagg aagttcaacg ccacagagag ggtcctgcag	120
ggtctgctta gtcccatatt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ctgaccttgc tcagacctga gaagcaggag gcagccactg gagtggacac catctgtacc	240
caccgcgttg atcccatcgg acctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca	420
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca	468
<210> 133 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 133 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan	60
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag	120
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga	180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc Page 130	240

caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360
gtcaatggtt tcacccatcg gagctttggg ctcaccacca gcactccttg gacttccaca	420
gttgaccttg gaacctcagg gactccatcc cccgtcccca gccccaca	468
<210> 134 <211> 468 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(468) <223> All N's = any nucleotide	
<400> 134 actgctggcc ctctcctggt gccattcacc ctaaacttca ccatcaccaa cctgcagtat	60
gaggaggaca tgcatcgccc tggctccagg aagttcaaca ccacggagag ggtccttcag	120
ggtctgctta cgcccttgtt caggaacacc agtgtcagct ctctgtactc tggttgcaga	180
ctgaccttgc tcaggcctga gaaggatggg gcagccacca gagtggatgc tgtctgcacc	240
catcgtcctg accccaaaag ccctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca	420
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca	468
<210> 135 <211> 465 <212> DNA <213> Homo sapiens	
<220> <221> misc_feature <222> (1)(465) <223> All N's = any nucleotide	
<400> 135 ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan	60
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag	120
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga	180
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc	240
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc	300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat	360

```
09-965738substitute.ST25.txt
                                                                       420
gtcaatggtt tcacccattg gatccctgtg cccaccagca gcactcctgg gacctccaca
                                                                       465
gtggaccttg ggtcagggac tccatcctcc ctccccagcc ccaca
<210>
      136
<211>
       468
<212>
      DNA
<213>
      Homo sapiens
<220>
      misc_feature
<221>
      (1)...(468)
All N's = any nucleotide
<222>
<223>
<400> 136
actgctggcc ctctcctggt accattcacc ctcaacttca ccatcaccaa cctgcagtat
                                                                        60
                                                                       120
ggggaggaca tgggtcaccc tggctccagg aagttcaaca ccacagagag ggtcctgcag
ggtctgcttg gtcccatatt caagaacacc agtgttggcc ctctgtactc tggctgcaga
                                                                       180
                                                                       240
ctgacctctc tcaggtccga gaaggatgga gcagccactg gagtggatgc Catctgcatc
catcatcttg accccaaaag ccctggactg nacagngagc ngctntactg ggagctnagc
                                                                       300
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat
                                                                       360
                                                                       420
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca
                                                                       468
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca
<210>
      137
<211>
       468
<212>
      DNA
<213> Homo sapiens
<220>
      misc_feature
<221>
      (1)..(468)
All N's = any nucleotide
<222>
<223>
<400> 137
ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan
                                                                        60
                                                                       120
gnggannaca tgcnncnccc nggntccagg aagttcaaca ccacngagng ngtnctgcag
ggtctgctnn nncccntntt caagaacncc agtgtnggcc ntctgtactc tggctgcaga
                                                                       180
                                                                       240
ctgacctnnc tcaggncnga gaagnatggn gcagccactg gantggatgc catctgcanc
                                                                       300
caccnncntn ancccaaaag ncctggactg nacagngagc ngctntactg ggagctnagc
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat
                                                                       360
gtcaatggtt tcacccatca gacctttgcg cccaacacca gcactcctgg gacctccaca
                                                                       420
                                                                       468
gtggaccttg ggacctcagg gactccatcc tccctccca gccctaca
```

```
<210>
      138
<211>
      468
<212>
      DNA
      Homo sapiens
<213>
<220>
      misc_feature
<221>
      (1)..(468)
<222>
<223> All N's = any nucleotide
<400> 138
                                                                      60
tctgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtac
gaggaggaca tgcatcaccc aggctccagg aagttcaaca ccacggagcg ggtcctgcag
                                                                      120
                                                                      180
qqtctqcttq qtcccatgtt caagaacacc agtgtcggcc ttctgtactc tggctgcaga
                                                                      240
ctgaccttgc tcaggcctga gaagaatggg gcagccacca gagtggatgc tgtctgcacc
                                                                      300
catcgtcctg accccaaaag ccctggactg nacagngagc ngctntactg ggagctnagc
canctgacca annncatcnn ngagctgggn ccctacaccc tggacaggna cagtctctat
                                                                      360
gtcaatggtt tcacccatcn ganctctgng cccaccacca gcactcctgg gacctccaca
                                                                      420
                                                                      468
gtgnacntng gnacctcngg gactccatcc tccntccccn gccncaca
<210>
      139
       468
<211>
<212>
      DNA
<213>
      Homo sapiens
<220>
<221> misc_feature
<222>
      (1)...(468)
<223> All N's = any nucleotide
<400>
      139
                                                                       60
ncnnctgncc ctctcctgnt nccnttcacc ntcaacttna ccatcaccaa cctgcantan
                                                                      120
qnqqannaca tgcnncnccc nggntccagg aagttcaaca ccacngagag ggttctgcag
                                                                      180
qqtctqctca aqcccttqtt caagagcacc agtgttggcc ctctgtattc tggctgcaga
ctgaccttgc tcaggcctga gaaggacgga gtagccacca gagtggacgc catctgcacc
                                                                      240
                                                                      300
caccgcctg accccaaaat ccctgggcta gacagacagc agctatactg ggagctgagc
                                                                      360
cagctgaccc acagcatcac tgagctggga ccctacaccc tggataggga cagtctctat
                                                                      420
gtcaatggtt tcacccagcg gagctctgtg cccaccacca gcactcctgg gactttcaca
                                                                      468
gtacagccgg aaacctctga gactccatca tccctccctg gccccaca
<210>
       140
       468
<211>
<212>
       DNA
```

<213>

Homo sapiens

09-965738substitute.ST25.txt <400> 140 gccactggcc ctgtcctgct gccattcacc ctcaatttta ccatcactaa cctgcagtat 60 120 gaggaggaca tgcatcgccc tggctccagg aagttcaaca ccacggagag ggtccttcag ggtctgctta tgcccttgtt caagaacacc agtgtcagct ctctgtactc tggttgcaga 180 240 ctgaccttgc tcaggcctga gaaggatggg gcagccacca gagtggatgc tgtctgcacc 300 catcgtcctg accccaaaag ccctggactg gacagagagc ggctgtactg gaagctgagc cagctgaccc acggcatcac tgagctgggc ccctacaccc tggacaggca cagtctctat 360 420 gtcaatggtt tcacccatca gagctctatg acgaccacca gaactcctga tacctccaca atgcacctgg caacctcgag aactccagcc tccctgtctg gacctacg 468 <210> 141 <211> 468 <212> DNA Homo sapiens <213> <400> 141 60 accgccagcc ctctcctggt gctattcaca attaacttca ccatcactaa cctgcggtat 120 gaggagaaca tgcatcaccc tggctctaga aagtttaaca ccacggagag agtccttcag 180 ggtctgctca ggcctgtgtt caagaacacc agtgttggcc ctctgtactc tggctgcaga ctgaccttgc tcaggcccaa gaaggatggg gcagccacca aagtggatgc catctgcacc 240 300 taccgccctg atcccaaaag ccctggactg gacagagagc agctatactg ggagctgagc cagctaaccc acagcatcac tgagctgggc ccctacaccc tggacaggga cagtctctat 360 gtcaatggtt tcacacagcg gagctctgtg cccaccacta gcattcctgg gacccccaca 420 468 gtggacctgg gaacatctgg gactccagtt tctaaacctg gtccctcg <210> 142 468 <212> DNA <213> Homo sapiens <400> 142 gctgccagcc ctctcctggt gctattcact ctcaacttca ccatcaccaa cctgcggtat 60 gaggagaaca tgcagcaccc tggctccagg aagttcaaca ccacggagag ggtccttcag 120 ggcctgctca ggtccctgtt caagagcacc agtgttggcc ctctgtactc tggctgcaga 180

240300

360

420 468

ctgactttgc tcaggcctga aaaggatggg acagccactg gagtggatgc catctgcacc

caccacctg acccaaaag ccctaggctg gacagagagc agctgtattg ggagctgagc

cagctgaccc acaatatcac tgagctgggc cactatgccc tggacaacga cagcctcttt

gtcaatggtt tcactcatcg gagctctgtg tccaccacca gcactcctgg gacccccaca

gtgtatctgg gagcatctaa gactccagcc tcgatatttg gcccttca

09-	-905/30Substitute.5125.txt
<210> 143 <211> 399 <212> DNA <213> Homo sapiens	
<400> 143 gctgccagcc atctcctgat actattca	cc ctcaacttca ccatcactaa cctgcggtat 60
gaggagaaca tgtggcctgg ctccagga	ag ttcaacacta cagagagggt ccttcagggc 120
ctgctaaggc ccttgttcaa gaacacca	gt gttggccctc tgtactctgg ctccaggctg 180
accttgctca ggccagagaa agatgggg	aa gccaccggag tggatgccat ctgcacccac 240
cgccctgacc ccacaggccc tgggctgg	ac agagagcagc tgtatttgga gctgagccag 300
ctgacccaca gcatcactga gctgggcc	cc tacacactgg acagggacag tctctatgtc 360
aatggtttca cccatcggag ctctgtac	cc accaccagc 399
<210> 144 <211> 453 <212> DNA <213> Homo sapiens <400> 144	
	ca ctgaacttca ccatcaacaa cctgcgctac 60
atggcggaca tgggccaacc cggctccc	tc aagttcaaca tcacagacaa cgtcatgaag 120
cacctgctca gtcctttgtt ccagagga	gc agcctgggtg cacggtacac aggctgcagg 180
gtcatcgcac taaggtctgt gaagaacg	gt gctgagacac gggtggacct cctctgcacc 240
tacctgcagc ccctcagcgg cccaggtc	tg cctatcaagc aggtgttcca tgagctgagc 300
cagcagaccc atggcatcac ccggctgg	gc ccctactctc tggacaaaga cagcctctac 360
cttaacggtt acaatgaacc tggtctag	at gagcctccta caactcccaa gccagccacc 420
acattcctgc ctcctctgtc agaagcca	ca aca 453
<210> 145 <211> 465 <212> DNA <213> Homo sapiens <400> 145	
	ca ctcaacttca ccatctccaa tctccagtat 60
tcaccagata tgggcaaggg ctcagcta	ca ttcaactcca ccgagggggt ccttcagcac 120
ctgctcagac ccttgttcca gaagagca	gc atgggcccct tctacttggg ttgccaactg 180
atctccctca ggcctgagaa ggatgggg	ca gccactggtg tggacaccac ctgcacctac 240
caccctgacc ctgtgggccc cgggctgg	ac atacagcagc tttactggga gctgagtcag 300
ctgacccatg gtgtcaccca actgggct	tc tatgtcctgg acagggatag cctcttcatc 360

465

146 <210> 9799 <211>

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222>

(1)..(9799) Any "X" = any amino acid <223>

<400>

Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Ala Thr Glu Arg Glu Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg

Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu 50 60

Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr 65 70 75 80

His Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr 85 90 95

Trp Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr $100 \hspace{1cm} 105 \hspace{1cm} 110$

Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser 115 120 125

Ser Met Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly

Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Ala Ala Gly Pro 145 150 155 160

Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr 165 170

Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Page 136

Ser Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val 195 200 205 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 210 220 Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp 225 230 235 240 Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser 245 250 255 Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg 260 265 270Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr 275 280 285 Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr 290 295 300 Pro Ser Ser Leu Ser Ser Pro Thr Ile Met Ala Ala Gly Pro Leu Leu 305 310 315 320 Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu 325 330 335 Asp Met Gly His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 340 345 350 Leu Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser Val Gly Pro 355 360 365 Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys Asp Gly 370 375 380 Ala Ala Thr Gly Val Asp Ala Ile Cys Ile His His Leu Asp Pro Lys 385 390 395 400 Ser Pro Gly Leu Asn Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu 405 410 415 Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser 420 425 430

09-965738substitute.ST25.txt Leu Tyr Val Asn Gly Phe Thr His Arg Thr Ser Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe Ser Leu Pro Ser Pro Ala Thr Ala Gly Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Thr Leu 500 505 510 Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly 530 540 Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile 565 570 575 Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn 580 585 590 Gly Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ala Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile 630 640 Thr Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys
645 650 655 Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe 660 Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu 675 680 685

Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys 690 700 Thr His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu 705 710 715 720 Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro 725 730 735 Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln
740 745 750 Thr Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu 755 760 765 Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly 770 780 Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 820 825 830 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu 835 840 845 Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu 850 860 Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu 865 870 875 880 Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp 885 890 895 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro 900 905 910 Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly 915 920 925 Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val 930 935 940 Page 139

- Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 945 950 955 960
- Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 965 970 975
- Gln Gly Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu 980 985 990
- Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asn Gly Ala 995 1000 1005
- Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu Asp Pro Lys 1010 1020
- Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln 1025 1030 1035
- Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg 1040 1045 1050
- Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala 1055 1060 1065
- Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser 1070 1080
- Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val Pro Leu 1085 1090 1095
- Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr 1100 11105 1110
- Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr 1115 1120 1125
- Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser 1130 1135 1140
- Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg 1145 1150 1155
- Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr 1160 1165 1170
- His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Page 140

1175

Туг	тгр 1190	Gln	Leu	Ser	Gln	Met 1195	Thr	Asn	Gly	Ile	Lys 1200	Glu	Leu	Gly
Pro	Tyr 1205	Thr	Leu	Asp	Arg	Asn 1210	Ser	Leu	туг	val	Asn 1215	Gly	Phe	Thr
His	Arg 1220	Ser	Ser	Gly	Leu	Thr 1225	Thr	Ser	Thr	Pro	Trp 1230	Thr	Ser	Thr
val	Asp 1235	Leu	Gly	Thr	Ser	Gly 1240	Thr	Pro	Ser	Pro	val 1245	Pro	Ser	Pro
Thr	Thr 1250	Ala	Glу	Pro	Leu	Leu 1255	٧a٦	Pro	Phe	Thr	Leu 1260	Asn	Phe	Thr
Ile	Thr 1265	Asn	Leu	Gln	Tyr	Glu 1270	Glu	Asp	Met	His	Arg 1275	Pro	Gly	Ser
Arg	Lys 1280	Phe	Asn	Ala	Thr	Glu 1285	Arg	٧a٦	Leu	Gln	Gly 1290	Leu	Leu	Ser
Pro	Ile 1295	Phe	Lys	Asn	Ser	Ser 1300	val	Gly	Pro	Leu	Tyr 1305	Ser	Gly	Cys
Arg	Leu 1310	Thr	Ser	Leu	Arg	Pro 1315	Glu	Lys	Asp	Gly	Ala 1320	Ala	Thr	Gly
Met	Asp 1325	Ala	٧a٦	Cys	Leu	Tyr 1330	His	Pro	Asn	Pro	Lys 1335	Arg	Pro	Gly
Leu	Asp 1340	Arg	Glu	Gl'n	Leu	Tyr 1345	Trp	Glu	Leu	Ser	Gln 1350	Leu	Thr	His
Asn	Ile 1355	Thr	Glu	Leu	Gly	Pro 1360	Tyr	Ser	Leu	Asp	Arg 1365	Asp	Ser	Leu
Tyr	Val 1370	Asn	Gly	Phe	Thr	His 1375	Gln	Asn	Ser	val	Pro 1380	Thr	Thr	Ser
Thr	Pro 1385	Gly	Thr	Ser	Thr	val 1390	Tyr	Trp	Ala	Thr	Thr 1395	Gly	Thr	Pro
Ser	Ser 1400	Phe	Pro	Gly	His	Thr 1405	Glu	Pro	Gly	Pro	Leu 1410	Leu	Ile	Pro

09-965738substitute.ST25.txt Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn 1420 Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 1430 1435 1440 Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys 1460 1470 Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 1475 1480 1485 Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys Glu 1495 1500 1490 Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser 1505 1510 1515 1510 1505 Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp 1535 1540 1545 Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe 1605 1600 Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr 1615 1610 Leu Leu Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr 1625 Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg

- Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr 1665 1655 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn 1670 1680 Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu 1700 1710 Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr 1760 1765 1770 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Glu Ala 1775 1780 1785 Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile 1790 Gly Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa 1805 Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg 1820 Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa 1840 1845 Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser 1855
- Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr 1890 1885 Page 143

Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Ser Ala Gly Pro Leu

1870

Glu	Glu 1895	Asp	Met	His	His	Pro 1900	Gly	Ser	Arg	Lys	Phe 1905	Asn	Thr	Thr
Glu	Arg 1910	∨al	Leu	Gln	Glу	Leu 1915	Leu	Gly	Pro	Met	Phe 1920	Lys	Asn	Thr
Ser	val 1925	Gly	Leu	Leu	Tyr	Ser 1930	Gly	Cys	Arg	Leu	Thr 1935	Leu	Leu	Arg
Pro	Glu 1940	Lys	Asn	Gly	Ala	Ala 1945	Thr	Glу	Met	Asp	Ala 1950	Ile	Cys	Ser
His	Arg 1955	Leu	Asp	Pro	Lys	Ser 1960	Pro	Gly	Leu	Asp	Arg 1965	Glu	Gln	Leu
Tyr	Trp 1970	Glu	Leu	Ser	Gln	Leu 1975	Thr	His	Gly	Ile	Lys 1980	Glu	Leu	Gly
Pro	Tyr 1985	Thr	Leu	Asp	Arg	Asn 1990	Ser	Leu	туг	۷al	Asn 1995	Gly	Phe	Thr
ніѕ	Arg 2000	Ser	Ser	۷al	Ala	Pro 2005	Thr	Ser	Thr	Pro	Gly 2010	Thr	Ser	Thr
٧a٦	Asp 2015	Leu	Gly	Thr	Ser	G]y 2020	Thr	Pro	Ser	Ser	Leu 2025	Pro	Ser	Pro
Thr	Thr 2030	Ala	٧a٦	Pro	Leu	Leu 2035	٧a٦	Pro	Phe	Thr	Leu 2040	Asn	Phe	Thr
Ile	Thr 2045	Asn	Leu	Gln	Tyr	Gly 2050	Glu	Asp	Met	Arg	ніs 2055	Pro	Gly	Ser
Arg	Lys 2060	Phe	Asn	Thr	Thr	G]u 2065	Arg	val	Leu	Gln	Gly 2070	Leu	Leu	Gly
Pro	Leu 2075	Phe	Lys	Asn	Ser	Ser 2080	val	Gly	Pro	Leu	Tyr 2085	Ser	Gly	Cys
Arg	Leu 2090	Ile	Ser	Leu	Arg	Ser 2095	Glu	Lys	Asp	Gly	Ala 2100	Ala	Thr	Gly
val	Asp 2105	Ala	Ile	Cys	Thr	His 2110	His	Leu	Asn	Pro	Gln 2115	Ser	Pro	Gly
Leu	Asp	Arg	Glu	Gln	Leu	Tyr	Trp	Gln	Leu Page	Ser e 14	Gln 4	Met	Thr	Asn

Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 2135 2140 2145 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser 2150 2160 Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro 2185 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val 2210 2215 2220 Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys 2240 2245 2250 Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro 2255 2260 2265 Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu 2270 Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser 2285 2290 2295 2285 Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser 2300 2310 Ser Met Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu 2320 2325 2315 Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Cys Thr Ile Thr Asn

2350

09-965738substitute.ST25.txt Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe 2360 2370 Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu Lys Pro Leu Phe 2375 2380 2385 Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr 2390 2400 Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly Val Asp Ala 2405 2410 2415 Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg 2420 2425 2430 2420 Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu 2440 2445 2435 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn 2450 2460 Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly 2465 2470 2475 Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser Leu 2480 2490 Ser Ser Pro Thr Ile Met Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 2525 2530 2535 Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp 2555 2560 2565 Gly Ala Ala Thr Arg Val Asp Ala Ala Cys Thr Tyr Arg Pro Asp 2570 2580 Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu 2585 2590 2595

Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu 2610 2600 Asp Arg Val Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser 2620 Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala 2630 Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu 2665 His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu 2705 2715 Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile 2720 2725 2730 2**7**Ž0 Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu 2735 Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu 2750 2760 Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 2765 2770 2775 Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr 2780 2785 2790 Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Val Asp Met Arg His Pro 2830 Page 147

Gly	Ser 2840	Arg	Lys	Phe	Asn	Thr 2845	Thr	Glu	Arg	val	Leu 2850	G]n	Gly	Leu
Leu	Lys 2855	Pro	Leu	Phe	Lys	Ser 2860	Thr	Ser	val	Gly	Pro 2865	Leu	Tyr	Ser
Gly	Cys 2870	Arg	Leu	Thr	Leu	Leu 2875	Arg	Pro	Glu	Lys	Arg 2880	Gly	Ala	Αla
Thr	Gly 2885		Asp	Thr	Ile	Cys 2890	Thr	His	Arg	Leu	Asp 2895	Pro	Leu	Asn
Pro	G]y 2900	Leu	Asp	Arg	Glu	G1n 2905	Leu	Tyr	Тгр	Glu	Leu 2910	Ser	Lys	Leu
Thr	Arg 2915	Gly	Ile	Ile	Glu	Leu 2920	Gly	Pro	Tyr	Leu	Leu 2925	Asp	Arg	Glу
Ser	Leu 2930	Tyr	۷al	Asn	Gly	Phe 2935	Thr	His	Arg	Asn	Phe 2940	val	Pro	Ile
Thr	Ser 2945	Thr	Pro	Gly	Thr	Ser 2950	Thr	val	His	Leu	Gly 2955	Thr	Ser	Glu
Thr	Pro 2960	Ser	Ser	Leu	Pro	Arg 2965	Pro	Ile	val	Pro	G]y 2970	Pro	Leu	Leu
val	Pro 2975	Phe	Thr	Leu	Asn	Phe 2980	Thr	Ile	Thr	Asn	Leu 2985	Gln	Tyr	Glu
Glu	Ala 2990	Met	Arg	ніѕ	Pro	Gly 2995	Ser	Arg	Lys	Phe	Asn 3000	Thr	Thr	Glu
Arg	val 3005	Leu	Gln	Gly	Leu	Leu 3010	Arg	Pro	Leu	Phe	Lys 3015	Asn	Thr	Ser
Ile	G1y 3020	Pro	Leu	туг	Ser	Ser 3025	Cys	Arg	Leu	Thr	Leu 3030	Leu	Arg	Pro
Glu	Lys 3035	Asp	Lys	Ala	Ala	Thr 3040	Arg	۷a٦	Asp	Ala	11e 3045	Cys	Thr	His
His	Pro 3050	Asp	Pro	Gln	Ser	Pro 3055	Gly	Leu	Asn	Arg	G]u 3060	Gln	Leu	Tyr
Trp	Glu	Leu	Ser	Gln	Leu	Thr	нis	Gly	Ile Page	Thr e 148	Glu 3	Leu	Gly	Pro

Tyr	Thr 3080	Leu	Asp	Arg	Asp	Ser 3085	Leu	Tyr	val	Asp	G]y 3090	Phe	Thr	ніѕ
Trp	Ser 3095	Pro	Ile	Pro	Thr	Thr 3100	Ser	Thr	Pro	Glу	Thr 3105	Ser	Ile	Val
Asn	Leu 3110	Gly	Thr	Ser	Gly	Ile 3115	Pro	Pro	Ser	Leu	Pro 3120	Glu	Thr	Thr
Xaa	xaa 3125	Xaa	Pro	Leu	Leu	Xaa 3130	Pro	Phe	Thr	Leu	Asn 3135	Phe	Thr	Ile
Thr	Asn 3140	Leu	Xaa	Tyr	Glu	Glu 3145	Xaa	Met	Xaa	Xaa	Pro 3150	Gly	Ser	Arg
Lys	Phe 3155	Asn	Thr	Thr	Glu	Arg 3160	val	Leu	Gln	Glу	Leu 3165	Leu	Lys	Pro
Leu	Phe 3170	Arg	Asn	Ser	Ser	Leu 3175	Glu	Tyr	Leu	туг	Ser 3180	Gly	Cys	Arg
Leu	Ala 3185	Ser	Leu	Arg	Pro	Glu 3190	Lys	Asp	Ser	Ser	Ala 3195	Met	Ala	val
Asp	Ala 3200	Ile	Cys	Thr	His	Arg 3205	Pro	Asp	Pro	Glu	Asp 3210	Leu	Gly	Leu
Asp	Arg 3215	Glu	Arg	Leu	Tyr	Trp 3220	Glu	Leu	Ser	Asn	Leu 3225	⊤hr	Asn	Gly
Ile	G]n 3230	Glu	Leu	Gly	Pro	Tyr 3235	Thr	Leu	Asp	Arg	Asn 3240	Ser	Leu	Tyr
val	Asn 3245	Gly	Phe	Thr	His	Arg 3250	Ser	Ser	Phe	Leu	Thr 3255	Thr	Ser	Thr
Pro	Trp 3260	Thr	Ser	Thr	val	Asp 3265	Leu	Gly	Thr	Ser	Gly 3270	Thr	Pro	Ser
Pro	va1 3275	Pro	Ser	Pro	Thr	Thr 3280	Ala	Gly	Pro	Leu	Leu 3285	Va]	Pro	Phe
Thr	Leu 3290	Asn	Phe	Thr	Ile	Thr 3295	Asn	Leu	Gln	Tyr	Glu 3300	Glu	Asp	Met

09-965738substitute_ST25.txt His Arg Pro Gly Ser Arg Arg Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val Asp 3350 3360 Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 3365 3370 3375 3365 Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu 3390 3380 Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser 3395 3400 val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala 3410 Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val 3425 3430 3435 Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asp Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg_ Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu 3485 3490 3495 Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile 3500 3510 35ŎO Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu 3515 Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu 3530 3540 3530 Page 150

Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 3545 3550 3555 Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr 3565 Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro 3580 Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro 3610 3615 Gly Ser Arg Lys Phe Ser Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala 3650 3660 Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser 3665 3670 3675 3665 Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu 3680 Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His 3695 Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr 3710 Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg 3725 3730 3735 Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu 3740 3750 3740 Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 3770 3780 Page 151

Arg	val 3785	Leu	Gln	Gly	Leu	Leu 3790	Arg	Pro	val	Phe	Lys 3795	Asn	Thr	Ser
val	Gly 3800	Pro	Leu	Tyr	Ser	Gly 3805	Cys	Arg	Leu	Thr	Leu 3810	Leu	Arg	Pro
Lys	Lys 3815	Asp	Gly	Ala	Ala	Thr 3820	Lys	val	Asp	Аlа	Ile 3825	Cys	Thr	Tyr
Arg	Pro 3830	Asp	Pro	Lys	Ser	Pro 3835	Gly	Leu	Asp	Arg	Glu 3840	Gln	Leu	Tyr
Тгр	Glu 3845	Leu	Ser	Gln	Leu	Thr 3850	His	Ser	Ile	Thr	G]u 3855	Leu	Gly	Pro
Туг	Thr 3860	Gln	Asp	Arg	Asp	Ser 3865	Leu	Tyr	val	Asn	Gly 3870	Phe	Thr	His
Arg	ser 3875	Ser	٧a٦	Pro	Thr	Thr 3880	Ser	Ile	Pro	Gly	Thr 3885	Ser	Ala	val
His	Leu 3890	Glu	Thr	Ser	Gly	Thr 3895	Pro	Ala	Ser	Leu	Pro 3900	Gly	His	Thr
Ala	Pro 3905	Gly	Pro	Leu	Leu	Val 3910	Pro	Phe	Thr	Leu	Asn 3915	Phe	Thr	Ile
Thr	Asn 3920	Leu	Gln	Tyr	Glu	Glu 3925	Asp	Met	Arg	His	Pro 3930	Gly	ser	Arg
Lys	Phe 3935	Asn	Thr	Thr	Glu	Arg 3940	val	Leu	Gln	Gly	Leu 3945	Leu	Lys	Pro
Leu	Phe 3950	Lys	ser	Thr	Ser	Val 3955	Gly	Pro	Leu	туг	ser 3960	Gly	Cys	Arg
Leu	Thr 3965	Leu	Leu	Arg	Pro	Glu 3970	Lys	Arg	Gly	Ala	Ala 3975	Thr	Gly	val
Asp	Thr 3980	Ile	Cys	Thr	His	Arg 3985	Leu	Asp	Pro	Leu	Asn 3990	Pro	Gly	Leu
Asp	Arg 3995	Glu	Gln	Leu	Tyr	Trp 4000	Glu	Leu	Ser	Lys	Leu 4005	Thr	Arg	Gly
Ile	Ile	Glu	Leu	Gly	Pro	Туг	Leu	Leu	Asp Pag	Arg e 15	Gly 2	Ser	Leu	Tyr

Val Asn Gly Phe Thr His Arg Thr Ser Val Pro Thr Thr Ser Thr 4030 4025 Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe 4040 4045 4050 Ser Leu Pro Ser Pro Ala Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met 4070 4075 4080 Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Thr Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp 4130 4140 Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu 4145 4150 4155 Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu 4160 Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Trp Ile Pro 4175 4180 4185 Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Leu Pro Ser Ser Pro Thr Thr Ala Gly Pro 4205 4210 4215 Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr

09-965738substitute.ST25.txt Thr Glu Arg Val Leu Gln Ser Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu 4310 4320 Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe 4325 4330 4335 Thr His Gln Thr Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser 4340 4345 Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe 4370 4380 Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr 4430 4440 Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro 4445 Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr 4470 4460 Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp_ Arg Xaa Ser 4475

Leu Tyr Val Asn Gly Phe Thr His Trp Ile Pro Val Pro Thr Ser 4500 4490 Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro 4505 Ser Ser Leu Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp 4545 Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 4550 4560 Leu Gln Ser Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys 4585 Asp Gly_ Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Val Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu 4615 4620 4610 Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 4635 4625 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr 4640 4650 Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala 4670 4680 Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn 4685 Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe 4725 4720 Page 155

Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala 4745 4750 4755 Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa 4775 4780 4785 Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn 4790 4795 4800 Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly 4815 4805 Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa 4820 Pro Xaa Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu 4840 Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa 4850 4860 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu Tyr 4885 Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu 4915 4920 4910 Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn 4925 4935 4930 Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg 4945 4950 4940 Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Page 156

Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser 4970 Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Thr Ala Gly Pro Leu 4990 Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe Asn Thr Thr 5015 5020 5025 Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Ile His His Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Arg Leu 5085 5080 Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly 5095 5090 Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr 5105 5110 His Arg Thr Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr 5120 5130 5120 Val Asp Leu Gly Thr Ser Gly Thr Pro Phe Ser Leu Pro Ser Pro Ala Thr Ala Gly Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr 5150 5160 Ile Thr Asn Leu Lys Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Thr Leu Leu Gly

09-965738substitute.ST25.txt Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly 5225 5235 Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa 5240 5250 Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu 5255 5260 5265 Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser 5280 5270 Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro 5285 xaa xaa xaa Pro xaa xaa Thr Xaa xaa xaa Pro Leu Leu Xaa Pro 5305 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa 5315 5320 5325 Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 5330 5340 Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys 5360 5370 Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro 5375 Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu 5395 5400 5390 Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr 5405 Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser 5420 5430 5420 Page 158

						U:	,- 50.	77 303	ubsi	Litu		. J . C	``	
Ser	Val 5435	Pro	Thr	Thr	Ser	Ile 5440	Pro	Gly	Thr	Ser	Ala 5445	val	His	Leu
Glu	Thr 5450	Thr	Gly	Thr	Pro	Ser 5455	Ser	Phe	Pro	Gly	ніs 5460	Thr	Glu	Pro
Gly	Pro 5465	Leu	Leu	Ile	Pro	Phe 5470	Thr	Phe	Asn	Phe	Thr 5475	Ile	Thr	Asn
Leu	Arg 5480	Tyr	Glu	Glu	Asn	Met 5485	Gln	His	Pro	Gly	Ser 5490	Arg	Lys	Phe
Asn	Thr 5495	Thr	Glu	Arg	val	Leu 5500	Gln	Gly	Leu	Leu	Thr 5505	Pro	Leu	Phe
Lys	Asn 5510	Thr	Ser	٧a٦	Gly	Pro 5515	Leu	Tyr	Ser	Gly	Cys 5520	Arg	Leu	Thr
Leu	Leu 5525	Arg	Pro	Glu	Lys	Gln 5530	Glu	Ala	Ala	Thr	Gly 5535	val	Asp	Thr
Ile	Cys 5540	Thr	His	Arg	val	Asp 5545	Pro	Ile	Gly	Pro	Gly 5550	Leu	Asp	Arg
Glu	Arg 5555	Leu	Tyr	Trp	Glu	Leu 5560	Ser	Gln	Leu	Thr	Asn 5565	Ser	Ile	Thr
Glu	Leu 5570	Gly	Pro	Tyr	Thr	Leu 5575	Asp	Arg	Asp	Ser	Leu 5580	Tyr	val	Asp
Gly	Phe 5585	Asn	Pro	Trp	Ser	Ser 5590	val	Pro	Thr	Thr	Ser 5595	Thr	Pro	Gly
Thr	Ser 5600	Thr	val	нis	Leu	Ala 5605	Thr	Ser	Gly	Thr	Pro 5610	Ser	Pro	Leu
Pro	Gly 5615	His	Thr	Ala	Pro	val 5620	Pro	Leu	Leu	Ile	Pro 5625	Phe	Thr	Leu
Asn	Phe 5630	Thr	Ile	Thr	Asp	Leu 5635	нis	Tyr	Glu	Glu	Asn 5640	Met	Gln	His
Pro	Gly 5645	Ser	Arg	Lys	Phe	Asn 5650	Thr	Thr	Glu	Arg	val 5655	Leu	Gln	Gly
Leu	Leu 5660	Lys	Pro	Leu	Phe	Lys 5665	Ser	Thr		Val e 159	5670	Pro	Leu	Tyr

Ser	G]y 5675	Cys	Arg	Leu	Thr	Leu 5680	Leu	Arg	Pro	Glu	Lys 5685	His	Gly	Ala
Ala	Thr 5690	Gly	val	Asp	Аlа	Ile 5695	Cys	Thr	Leu	Arg	Leu 5700	Asp	Pro	Thr
Gly	Pro 5705	Gly	Leu	Asp	Arg	Glu 5710	Arg	Leu	Tyr	Тгр	Glu 5715	Leu	Ser	Gln
Leu	Thr 5720	Asn	Ser	Ile	Thr	G1u 5725	Leu	Gly	Pro	Tyr	Thr 5730	Leu	Asp	Arg
Asp	ser 5735	Leu	Tyr	val	Asn	G1y 5740	Phe	Asn	Pro	Тгр	Ser 5745	Ser	val	Pro
Thr	Thr 5750	Ser	Thr	Pro	Gly	Thr 5755	Ser	Thr	val	His	Leu 5760	Ala	Thr	Ser
Gly	Thr 5765	Pro	Ser	Ser	Leu	Pro 5770	Gly	His	Thr	Thr	Ala 5775	Gly	Pro	Leu
Leu	val 5780	Pro	Phe	Thr	Leu	Asn 5785	Phe	Thr	Ile	Thr	Asn 5790	Leu	Lys	Tyr
Glu	Glu 5795	Asp	Met	нis	Cys	Pro 5800	Gly	Ser	Arg	Lys	Phe 5805	Asn	Thr	Thr
Glu	Arg 5810		Leu	Gln	Ser	Leu 5815	His	Gly	Pro	Met	Phe 5820	Lys	Asn	Thr
Ser	Val 5825	Gly	Pro	Leu	Tyr	Ser 5830	Gly	Cys	Arg	Leu	Thr 5835	Leu	Leu	Arg
Ser	Glu 5840	Lys	Asp	Gly	Ala	Ala 5845	Thr	Gly	٧a٦	Asp	Ala 5850	Ile	Cys	Thr
His	Arg 5855	Leu	Asp	Pro	Lys	Ser 5860	Pro	Gly	Leu	Asp	Arg 5865	Glu	Xaa	Leu
Tyr	Trp 5870	Glu	Leu	Ser	xaa	Leu 5875	Thr	Xaa	xaa	Ile	Xaa 5880	Glu	Leu	Gly
Pro	Tyr 5885	Xaa	Leu	Asp	Arg	Xaa 5890	Ser	Leu	Tyr	val	Asn 5895	Gly	Phe	Xaa
xaa	xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Thr	Ser	Thr Page	Pro e 160	Gly O	Thr	Ser	xaa

	5900					5905	9-965	57389	subst	titu	5910	25.t	ĸt	
Val	Xaa 5915	Leu	Xaa	Thr	Ser	Gly 5920	Thr	Pro	xaa	xaa	Xaa 5925	Pro	Xaa	Xaa
Thr	Xaa 5930	Xaa	xaa	Pro	Leu	Leu 5935	xaa	Pro	Phe	Thr	Leu 5940	Asn	Phe	Thr
Ile	Thr 5945	Asn	Leu	Xaa	Туг	G]u 5950	Glu	Xaa	Met	xaa	xaa 5955	Pro	Gly	Ser
Arg	Lys 5960	Phe	Asn	Thr	Thr	Glu 5965	Arg	∨al	Leu	Gln	Gly 5970	Leu	Leu	Xaa
Pro	Xaa 5975	Phe	Lys	Xaa	Thr	Ser 5980	val	Gly	xaa	Leu	Tyr 5985	Ser	Gly	Cys
Arg	Leu 5990	Thr	Leu	Leu	Arg	Xaa 5995	Glu	Lys	xaa	xaa	Ala 6000	Ala	Thr	xaa
۷al	Asp 6005	xaa	xaa	Cys	xaa	Xaa 6010	Xaa	xaa	Asp	Pro	xaa 6015	Xaa	Pro	Gly
Leu	Asp 6020	Arg	Glu	xaa	Leu	Tyr 6025	Trp	Glu	Leu	Ser	xaa 6030	Leu	Thr	Asn
Ser	11e 6035	Thr	Glu	Leu	Gly	Pro 6040	Tyr	Thr	Leu	Asp	Arg 6045	Asp	Ser	Leu
Tyr	Val 6050	Asn	Gly	Phe	Thr	His 6055	Arg	Ser	Ser	Met	Pro 6060	Thr	Thr	Ser
Ile	Pro 6065	Gly	Thr	Ser	Ala	val 6070	His	Leu	Glu	Thr	Ser 6075	Gly	Thr	Pro
Ala	Ser 6080	Leu	Pro	Gly	нis	Thr 6085	Ala	Pro	Gly	Pro	Leu 6090	Leu	val	Pro
Phe	Thr 6095	Leu	Asn	Phe	Thr	Ile 6100	Thr	Asn	Leu	Gln	туг 6105	Glu	Glu	Asp
Met	Arg 6110	His	Pro	Gly	Ser	Arg 6115	Lys	Phe	Asn	Thr	Thr 6120	Glu	Arg	val
Leu	Gln 6125	Gly	Leu	Leu	Lys	Pro 6130	Leu	Phe	Lys	Ser	Thr 6135	Ser	val	Gly

09-965738substitute.ST25.txt Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa 6185 6190 6195 Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa 6200 Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu 6220 6215 Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa 6230 Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe 6260 6270 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg 6320 6330 Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa 6345 6335 Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn 6360 6350 Gly Phe His Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly 6370 6365

Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu 6390 6380 Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu 6400 Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His 6415 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 6425 6430 6435 Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr 6440 6455 6450 ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa 6490 Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg 6510 6500 6505 Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa 6515 Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser 6540 6530 Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr 6560 6570 Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr 6575 6580 Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg 6610 Page 163

Xaa	G1u 6620	Lys	Xaa	xaa	Ala	Ala 6625	Thr	xaa	val	Asp	Xaa 6630	xaa	Cys	xaa
Xaa	xaa 6635	xaa	Asp	Pro	xaa	Xaa 6640	Pro	GТу	Leu	Asp	Arg 6645	Glu	xaa	Leu
Tyr	Trp 6650	Glu	Leu	Ser	Xaa	Leu 6655	Thr	Xaa	Xaa	Ile	Xaa 6660	Glu	Leu	Gly
Pro	Tyr 6665	Xaa	Leu	Asp	Arg	Xaa 6670	Ser	Leu	туг	val	Asn 6675	Gly	Phe	Thr
His	Gln 6680	Asn	Ser	val	Pro	Thr 6685	Thr	Ser	Thr	Pro	G]y 6690	Thr	Ser	Thr
٧a٦	Tyr 6695	Trp	Ala	Thr	Thr	Gly 6700	Thr	Pro	Ser	Ser	Phe 6705	Pro	Gly	His
Thr	Glu 6710	Pro	Gly	Pro	Leu	Leu 67 1 5	Ile	Pro	Phe	Thr	Phe 6720	Asn	Phe	Thr
Ile	Thr 6725	Asn	Leu	His	Tyr	Glu 6730	Glu	Asn	Met	Gln	ніs 6735	Pro	Gly	Ser
Arg	Lys 6740	Phe	Asn	Thr	Thr	Glu 6745	Arg	val	Leu	Gln	Gly 6750	Leu	Leu	Thr
Pro	Leu 6755	Phe	Lys	Asn	Thr	ser 6760	val	Gly	Pro	Leu	Tyr 6765	Ser	Gly	Cys
Arg	Leu 6770	Thr	Leu	Leu	Arg	Pro 6775	Glu	Lys	Gln	Glu	Ala 6780	Ala	Thr	Gly
٧al	Asp 6785	Thr	Ile	Cys	Thr	ніs 6790	Arg	٧a٦	Asp	Pro	Ile 6795	Gly	Pro	Gly
Leu	Asp 6800	Arg	Glu	xaa	Leu	Tyr 6805	Trp	Glu	Leu	Ser	xaa 6810	Leu	Thr	Xaa
Xaa	Ile 6815	Xaa	Glu	Leu	Gly	Pro 6820	Tyr	xaa	Leu	Asp	Arg 6825	Xaa	Ser	Leu
Tyr	va1 6830	Asn	Gly	Phe	Xaa	Xaa 6835	Xaa	Xaa	Xaa	Xaa	Xaa 6840	Xaa	Thr	Ser
Thr	Pro	Ģ٦y	Thr	Ser	Xaa	۷a٦	Xaa	Leu	Xaa Page	Thr e 16	Ser 4	Gly	Thr	Pro

Xaa Xaa 6860	Xaa	Pro	Xaa	Xaa	Thr 6865	Xaa	Xaa	xaa	Pro	Leu 6870	Leu	Xaa	Pro
Phe Thr 6875	Leu	Asn	Phe	Thr	Ile 6880	Thr	Asn	Leu	xaa	туг 6885	Glu	Glu	Xaa
Met Xaa 6890	Xaa	Pro	Gly	Ser	Arg 6895	Lys	Phe	Asn	Thr	Thr 6900	Glu	Arg	Val
Leu Gln 6905	Gly	Leu	Leu	xaa	Pro 6910	Xaa	Phe	Lys	Xaa	Thr 6915	Ser	val	Gly
Xaa Leu 6920		Ser	Gly	Cys	Arg 6925	Leu	Thr	Leu	Leu	Arg 6930	xaa	Glu	Lys
Xaa Xaa 6935	Ala	Аlа	Thr	xaa	Val 6940	Asp	Xaa	xaa	Cys	xaa 6945	xaa	xaa	Xaa
Asp Pro 6950	xaa	Xaa	Pro	Gly	Leu 6955	Asp	Arg	Glu	xaa	Leu 6960	Tyr	Тгр	Glu
Leu Ser 6965	xaa	Leu	Thr	Xaa	xaa 6970	Ile	xaa	Glu	Leu	Gly 6975	Pro	Tyr	Xaa
Leu Asp 6980		xaa	Ser	Leu	туг 6985	val	Asn	Gly	Phe	Thr 6990	His	Arg	Ser
Ser Val 6995	Pro	Thr	Thr	Ser	Ser 7000	Pro	Gly	Thr	Ser	Thr 7005	٧a٦	His	Leu
Ala Thr 7010	Ser	Gly	Thr	Pro	Ser 7015	Ser	Leu	Pro	Glу	His 7020	Thr	Ala	Pro
Val Pro 7025	Leu	Leu	Ile	Pro	Phe 7030	Thr	Leu	Asn	Phe	Thr 7035	Ile	Thr	Asn
Leu His 7040	туг	Glu	Glu	Asn	Met 7045	Gln	His	Pro	Gly	Ser 7050	Arg	Lys	Phe
Asn Thr 7055	Thr	Glu	Arg	val	Leu 7060	Gln	Gly	Leu	Leu	Lys 7065	Pro	Leu	Phe
Lys Ser 7070	Thr	Ser	٧a٦	Gly	Pro 7075	Leu	Tyr	Ser	Gly	Cys 7080	Arg	Leu	Thr

09-965738substitute_ST25.txt Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn 7130 Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly 7145 7150 7155 7145 Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa 7160 Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 7205 7210 7215 Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr 7220 7230 Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala 7235 7240 7245 Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg 7280 Xaa Ser Leu Tyr Val Asn Gly Phe Thr His Arg Thr Ser Val Pro 7295 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser 7310 Page 166

Page 167

Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr 7340 7350 Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser 7370 7375 7380 Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg 7385 7390 7395 Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu 7415 7420 7425 Tyr Cys Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly 7430 7440 Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr 7445 7450 7455 His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr 7460 7470 7460 Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His 7475 7480 7485 Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr 7495 7490 Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser 7505 7510 7515 Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa 7520 7530 Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa

Val Asp Xaa Xaa Cys Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly 7565 7570 7575 Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa 7580 7590 Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu 7595 7600 7605 Tyr Val Asn Gly Phe Thr His Trp Ser Ser Gly Leu Thr Thr Ser 7610 7620 Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro 7625 7630 Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro 7645 7650 7640 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 7655 Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val 7670 7680 Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Thr Ser Val Gly 7685 7690 7695 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 7700 7710 Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val 7715 7720 7725 Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu 7730 7740 Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa 7745 7750 7755 Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa 7760 7770 Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu 7775 7780 7785 Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Page 168

Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn 7805 7810 Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe 7825 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe 7840 Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr 7850 7860 Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa 7865 7870 7875 Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg 7880 7885 7890 Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn 7915 7920 7910 Gly Phe Thr His Arg Ser Phe Gly Leu Thr Thr Ser Thr Pro Trp 7925 Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val 7940 7945 7950 7940 Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu 7955 7960 7965 7955 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 7985 7990 7995 Leu Leu Thr Pro Leu Phe Arg Asn Thr Ser Val Ser Ser Leu Tyr 8000 8010 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 8015 8020 8025 8020

09-965738substitute.ST25.txt Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys 8035 8040 Ser Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa 8075 8080 Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser 8095 8100 8090 Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu 8105 8115 Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr 8130 8120 8125 Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr 8160 Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu 8200 Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Thr 8235 8225 8230 His Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr 8245 8240 Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr 8265 8255 8260

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile 8275 8270 8280 Thr Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg 8285 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro 8300 8310 Ile Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg 8315 8320 8325 Leu Thr Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val 8335 Asp Ala Ile Cys Ile His His Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa 8365 Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr 8380 8385 Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr 8400 8390 Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa 8415 8405 8410 Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe 8430 8420 Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met 8435 Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 8450 8460 Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa 8465 Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp 8500 8505 Page 171

Pro	xaa 8510	Xaa	Pro	Gly	Leu	Asp 8515	Arg	Glu	Xaa	Leu	Tyr 8520	Тгр	Glu	Leu
Ser	Xaa 8525	Leu	Thr	Xaa	Xaa	11e 8530	Xaa	Glu	Leu	Gly	Pro 8535	Tyr	Xaa	Leu
Asp	Arg 8540	xaa	Ser	Leu	Tyr	Val 8545	Asn	Gly	Phe	Thr	ніs 8550	Gln	Thr	Phe
Ala	Pro 8555	Asn	Thr	Ser	Thr	Pro 8560	Gly	Thr	Ser	Thr	va1 8565	Asp	Leu	Gly
Thr	Ser 8570	Gly	Thr	Pro	Ser	Ser 8575	Leu	Pro	Ser	Pro	Thr 8580	Ser	Ala	Gly
Pro	Leu 8585	Leu	٧al	Pro	Phe	Thr 8590	Leu	Asn	Phe	Thr	Ile 8595	Thr	Asn	Leu
Gln	Tyr 8600	Glu	Glu	Asp	Met	ніs 8605	His	Pro	Gly	Ser	Arg 8610	Lys	Phe	Asn
Thr	Thr 8615	Glu	Arg	∨al	Leu	G]n 8620	Glу	Leu	Leu	Gly	Pro 8625	Met	Phe	Lys
Asn	Thr 8630	Ser	val	Gly	Leu	Leu 8635	туг	Ser	Gly	Cys	Arg 8640	Leu	Thr	Leu
Leu	Arg 8645		Glu	Lys	Asn	Gly 8650	Ala	Ala	Thr	Arg	val 8655	Asp	Ala	val
Cys	Thr 8660	ніѕ	Arg	Pro	Asp	Pro 8665	Lys	Ser	Pro	Gly	Leu 8670	Asp	Arg	Glu
xaa	Leu 8675	Tyr	тгр	Glu	Leu	ser 8680	xaa	Leu	Thr	Xaa	Xaa 8685	Ile	Xaa	Glu
Leu	Gly 8690	Pro	Tyr	Xaa	Leu	Asp 8695	Arg	Xaa	Ser	Leu	Tyr 8700	٧a٦	Asn	Gly
Phe	Xaa 8705	Xaa	Xaa	xaa	Xaa	Xaa 8710	Xaa	xaa	Thr	Ser	Thr 8715	Pro	Gly	Thr
Ser	xaa 8720	val	xaa	Leu	Xaa	Thr 8725	Ser	Gly	Thr	Pro	Xaa 8730	Xaa	Xaa	Pro
xaa	xaa	Thr	Xaa	Xaa	Xaa	Pro	Leu	Leu	Xaa Pag	Pro e 17	Phe 2	Thr	Leu	Asn

Phe Thr 8750		Thr	Asn	Leu	Xaa 8755	Tyr	Glu	Glu	Xaa	Met 8760	xaa	xaa	Pro
Gly Ser 8765		Lys	Phe	Asn	Thr 8770	Thr	Glu	Arg	٧al	Leu 8775	Gln	Gly	Leu
Leu Lys 8780	Pro	Leu	Phe	Lys	Ser 8785	Thr	Ser	٧al	Gly	Pro 8790	Leu	Tyr	Ser
Gly Cys 8795	Arg	Leu	Thr	Leu	Leu 8800	Arg	Pro	Glu	Lys	Asp 8805	Gly	val	Ala
Thr Arg 8810		Asp	Ala	Ile	Cys 8815	Thr	His	Arg	Pro	Asp 8820	Pro	Lys	Ile
Pro Gly 8825		Asp	Arg	Gln	G1n 8830	Leu	Tyr	Trp	Glu	Leu 8835	Ser	Gln	Leu
Thr His 8840		Ile	Thr	Glu	Leu 8845	Gly	Pro	Туг	Thr	Leu 8850	Asp	Arg	Asp
Ser Leu 8855		٧a٦	Asn	Gly	Phe 8860	Thr	Gln	Arg	Ser	Ser 8865	val	Pro	Thr
Thr Ser 8870	Thr	Pro	Gly	Thr	Phe 8875	Thr	٧a٦	Gln	Pro	Glu 8880	Thr	Ser	Glu
Thr Pro 8885		Ser	Leu	Pro	Gly 8890	Pro	Thr	Ala	Thr	Gly 8895	Pro	val	Leu
Leu Pro 8900		Thr	Leu	Asn	Phe 8905	Thr	Ile	Thr	Asn	Leu 8910	Gln	Tyr	Glu
Glu ['] Asp 8915	Met	нis	Arg	Pro	Gly 8920	Ser	Arg	Lys	Phe	Asn 8925	Thr	Thr	Glu
Arg Val 8930		Gln	Gly	Leu	Leu 8935	Met	Pro	Leu	Phe	Lys 8940	Asn	Thr	Ser
Val Ser 8945		Leu	Tyr	Ser	Gly 8950	Cys	Arg	Leu	Thr	Leu 8955	Leu	Arg	Pro
Glu Lys 8960		Gly	Ala	Ala	Thr 8965	Arg	Val	Asp	Ala	Val 8970	Cys	Thr	His

09-965738substitute.ST25.txt Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu_ Arg Leu Tyr 8980 8985 Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro 9000 Tyr Thr Leu Asp Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Arg Thr Pro Asp Thr Ser Thr Met 9020 9025 9030 His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr 9045 9040 9035 Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile 9055 9050 Thr Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg 9070 9065 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Pro 9080 9085 9090 Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg 9095 9100 9105 Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val 9110 Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr 9165 Asn Val Gly Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Val 9170 9175 9180 Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Val 9185 Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val Leu Phe

Thr Leu Asn Gly Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met 9225 9215 Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 9230 9240 Gln Gly Leu Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp 9260 9270 Gly Thr Ala Thr Gly Val Asp Ala Ile Cys Thr His His Pro Asp Pro Lys Ser Pro Arg Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly His Tyr Ala Leu Asp Asn Asp Ser Leu Phe Val Asn Gly Phe Thr His Arg Ser Ser val Ser Thr Thr Ser Thr Pro Gly Thr Pro Thr Val Tyr Leu Gly 9345 9335 9340 Ala Ser Lys Thr Pro Ala Ser Ile Phe Gly Pro Ser Ala Ala Ser 9360 9350 His Leu Leu Ile Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu 9375 9365 Arg Tyr Glu Glu Asn Met Trp Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn 9395 9400 9405 9405 Thr Ser Val Gly Pro Leu Tyr Ser Gly Ser Arg Leu Thr Leu Leu 9410 Arg Pro Glu Lys Asp Gly Glu Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Gln 9445 Page 175

Leu	Tyr 9455	Leu	Glu	Leu	Ser	Gln 9460		Thr	His	Ser	11e 9465	Thr	Glu	Leu
Gly	Pro 9470	Tyr	Thr	Leu	Asp	Arg 9475	Asp	Ser	Leu	туг	va1 9480	Asn	Gly	Phe
Thr	ніs 9485	Arg	Ser	Ser	val	Pro 9490	Thr	Thr	Ser	Thr	Gly 9495	۷al	۷al	Ser
Glu	G]u 9500	Pro	Phe	Thr	Leu	Asn 9505	Phe	Thr	Ile	Asn	Asn 9510	Leu	Arg	Tyr
Met	Ala 9515	Asp	Met	Gly	Gln	Pro 9520	Gly	Ser	Leu	Lys	Phe 9525	Asn	Ile	Thr
Asp	Asn 9530	٧a٦	Met	Lys	His	Leu 9535	Leu	Ser	Pro	Leu	Phe 9540	Gln	Arg	Ser
Ser	Leu 9545	Gly	Ala	Arg	Tyr	Thr 9550	Gly	Cys	Arg	val	Ile 9555	Ala	Leu	Arg
Ser	Val 9560	Lys	Asn	Gly	Ala	Glu 9565	Thr	Arg	٧a٦	Asp	Leu 9570	Leu	Cys	Thr
Tyr	Leu 9575	Gln	Pro	Leu	Ser	Gly 9580		Gly	Leu	Pro	Ile 9585	Lys	Gln	val
Phe	ніs 9590	Glu	Leu	Ser	Gln	G]n 9595	Thr	His	Glý	Ile	Thr 9600	Arg	Leu	Gly
Pro	Tyr 9605	Ser	Leu	Asp	Lys	Asp 9610	Ser	Leu	Tyr	Leu	Asn 9615	Gly	Tyr	Asn
Glu	Pro 9620	Gly	Leu	Asp	Glu	Pro 9625	Pro	Thr	Thr	Pro	Lys 9630	Pro	Ala	Thr
Thr	Phe 9635	Leu	Pro	Pro	Leu	Ser 9640	Glu	Ala	Thr	Thr	Ala 9645	Met	Gly	Tyr
His	Leu 9650		Thr	Leu	Thr	Leu 9655	Asn	Phe	Thr	Ile	ser 9660	Asn	Leu	Gln
Туг	Ser 9665	Pro	Asp	Met	Gly	Lys 9670	Gly	Ser	Ala	Thr	Phe 9675	Asn	Ser	Thr
Glu	Gly	۷al	Leu	Gln	His	Leu	Leu	Arg	Pro Page	Leu e 17	Phe 6	Gln	Lys	Ser

Ser Met Gly Pro Phe Tyr Leu Gly Cys Gln Leu Ile Ser Leu Arg

Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr Thr Cys Thr 9710 9720

Tyr His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile Gln Gln Leu 9725 9730 9735

Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln Leu Gly 9740 9745 9750

Phe Tyr Val Leu Asp Arg Asp Ser Leu Phe Ile Asn Gly Tyr Ala 9755 9760 9765

Pro Gln Asn Leu Ser Ile Arg Gly Glu Tyr Gln Ile Asn Phe His 9770 9780

Ile Val Asn Trp Asn Leu Ser Asn Pro Asp Pro Thr Ser Ser Glu 9785 9790 9795

Tyr

<210> 147 <211> 1422 <212> DNA

9680

9695

<213> Homo sapiens

<400> 147

gccatggggt accacctgaa gaccctcaca ctcaacttca ccatctccaa tctccagtat 60 tcaccagata tgggcaaggg ctcagctaca ttcaactcca ccgagggggt ccttcagcac 120 180 ctgctcagac ccttgttcca gaagagcagc atgggcccct tctacttggg ttgccaactg atctccctca ggcctgagaa ggatggggca gccactggtg tggacaccac ctgcacctac 240 300 caccctgacc ctgtgggccc cgggctggac atacagcagc tttactggga gctgagtcag 360 ctgacccatg gtgtcaccca actgggcttc tatgtcctgg acagggatag cctcttcatc aatggctatg caccccagaa tttatcaatc cggggcgagt accagataaa tttccacatt 420 480 gtcaactgga acctcagtaa tccagacccc acatcctcag agtacatcac cctgctgagg 540 gacatccagg acaaggtcac cacactctac aaaggcagtc aactacatga cacattccgc ttctgcctgg tcaccaactt gacgatggac tccgtgttgg tcactgtcaa ggcattgttc 600 tcctccaatt tggaccccag cctggtggag caagtctttc tagataagac cctgaatgcc 660

tcattccatt	ggctgggctc	cacctaccag	ttggtggaca	tccatgtgac	agaaatggag	720
tcatcagttt	atcaaccaac	aagcagctcc	agcacccagc	acttctacct	gaatttcacc	780
atcaccaacc	taccatattc	ccaggacaaa	gcccagccag	gcaccaccaa	ttaccagagg	840
aacaaaagga	atattgagga	tgcgctcaac	caactcttcc	gaaacagcag	catcaagagt	900
tatttttctg	actgtcaagt	ttcaacattc	aggtctgtcc	ccaacaggca	ccacaccggg	960
gtggactccc	tgtgtaactt	ctcgccactg	gctcggagag	tagacagagt	tgccatctat	1020
gaggaatttc	tgcggatgac	ccggaatggt	acccagctgc	agaacttcac	cctggacagg	1080
agcagtgtcc	ttgtggatgg	gtattctccc	aacagaaatg	agcccttaac	tgggaattct	1140
gaccttccct	tctgggctgt	catcctcatc	ggcttggcag	gactcctggg	actcatcaca	1200
tgcctgatct	gcggtgtcct	ggtgaccacc	cgccggcgga	agaaggaagg	agaatacaac	1260
gtccagcaac	agtgcccagg	ctactaccag	tcacacctag	acctggagga	tctgcaatga	1320
ctggaacttg	ccggtgcctg	gggtgccttt	ccccagcca	gggtccaaag	aagcttggct	1380
ggggcagaaa	taaaccatat	tggtcggaaa	aaaaaaaaa	aa		1422

<210> 148

<211> 439

<212> PRT

<213> Homo sapiens

<400> 148

Ala Met Gly Tyr His Leu Lys Thr Leu Thr Leu Asn Phe Thr Ile Ser 1 10 15

Asn Leu Gln Tyr Ser Pro Asp Met Gly Lys Gly Ser Ala Thr Phe Asn 20 25 30

Ser Thr Glu Gly Val Leu Gln His Leu Leu Arg Pro Leu Phe Gln Lys 35 40 45

Ser Ser Met Gly Pro Phe Tyr Leu Gly Cys Gln Leu Ile Ser Leu Arg 50 60

Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr Thr Cys Thr Tyr 65 70 75 80

His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile Gln Gln Leu Tyr Trp 85 90 95

Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln Leu Gly Phe Tyr Val 100 105 110

Leu Asp Arg Asp Ser Leu Phe Ile Asn Gly Tyr Ala Pro Gln Asn Leu Page 178

Ser Ile Arg Gly Glu Tyr Gln Ile Asn Phe His Ile Val Asn Trp Asn 130 135 140 Leu Ser Asn Pro Asp Pro Thr Ser Ser Glu Tyr Ile Thr Leu Leu Arg 145 150 155 160 Asp Ile Gln Asp Lys Val Thr Thr Leu Tyr Lys Gly Ser Gln Leu His 165 170 175 Asp Thr Phe Arg Phe Cys Leu Val Thr Asn Leu Thr Met Asp Ser Val 180 185 190 Leu Val Thr Val Lys Ala Leu Phe Ser Ser Asn Leu Asp Pro Ser Leu 195 200 205 Val Glu Gln Val Phe Leu Asp Lys Thr Leu Asn Ala Ser Phe His Trp 210 215 220 Leu Gly Ser Thr Tyr Gln Leu Val Asp Ile His Val Thr Glu Met Glu 225 230 235 240 Ser Ser Val Tyr Gln Pro Thr Ser Ser Ser Ser Thr Gln His Phe Tyr 245 250 255 Leu Asn Phe Thr Ile Thr Asn Leu Pro Tyr Ser Gln Asp Lys Ala Gln 260 265 270 260 Pro Gly Thr Thr Asn Tyr Gln Arg Asn Lys Arg Asn Ile Glu Asp Ala 275 280 285 Leu Asn Gln Leu Phe Arg Asn Ser Ser Ile Lys Ser Tyr Phe Ser Asp 290 295 300 Cys Gln Val Ser Thr Phe Arg Ser Val Pro Asn Arg His His Thr Gly 305 310 315 Val Asp Ser Leu Cys Asn Phe Ser Pro Leu Ala Arg Arg Val Asp Arg 325 330 335 Val Ala Ile Tyr Glu Glu Phe Leu Arg Met Thr Arg Asn Gly Thr Gln 340 350 Leu Gln Asn Phe Thr Leu Asp Arg Ser Ser Val Leu Val Asp Gly Tyr 355 360 365

09-965738substitute.ST25.txt Ser Pro Asn Arg Asn Glu Pro Leu Thr Gly Asn Ser Asp Leu Pro Phe 370 380 Trp Ala Val Ile Leu Ile Gly Leu Ala Gly Leu Leu Gly Leu Ile Thr 385 390 395 400 Cys Leu Ile Cys Gly Val Leu Val Thr Thr Arg Arg Arg Lys Lys Glu 405 410 415 Gly Glu Tyr Asn Val Gln Gln Gln Cys Pro Gly Tyr Tyr Gln Ser His 420 430 Leu Asp Leu Glu Asp Leu Gln <210> 149 <211> 1799 <212> PRT <213> Homo sapiens <400> 149 Arg Thr Asp Gly Ile Met Glu His Ile Thr Lys Ile Pro Asn Glu Ala 1 10 15 Ala His Arg Gly Thr Ile Arg Pro Val Lys Gly Pro Gln Thr Ser Thr 20 25 30 Ser Pro Ala Ser Pro Lys Gly Leu His Thr Gly Gly Thr Lys Arg Met 35 40 45Glu Thr Thr Thr Ala Leu Lys Thr Thr Thr Ala Leu Lys Thr 50 60 Thr Ser Arg Ala Thr Leu Thr Thr Ser Val Tyr Thr Pro Thr Leu Gly 65 70 75 80 Thr Leu Thr Pro Leu Asn Ala Ser Arg Gln Met Ala Ser Thr Ile Leu 85 90 95 Thr Glu Met Met Ile Thr Thr Pro Tyr Val Phe Pro Asp Val Pro Glu 100 105 110 Thr Thr Ser Ser Leu Ala Thr Ser Leu Gly Ala Glu Thr Ser Thr Ala 115 120 125 Leu Pro Arg Thr Thr Pro Ser Val Leu Asn Arg Glu Ser Glu Thr Thr 130 140

Ala 145	Ser	Leu	Val	Ser	Arg 150	Ser							val		Gln 160
Thr	Leu	Asp	val	Ser 165	Ser	Ser	Glu	Pro	Asp 170	Thr	Thr	Ala	Ser	Trp 175	val
Ile	His	Pro	Ala 180	Glu	Thr	Ile	Pro	Thr 185	Val	Ser	Lys	Thr	Thr 190	Pro	Asn
Phe	Phe	ніs 195	Ser	Glu	Leu	Asp	Thr 200	٧al	Ser	Ser	Thr	Ala 205	Thr	Ser	His
Gly	Ala 210	Asp	val	Ser	Ser	Ala 215	Ile	Pro	Thr	Asn	11e 220	Ser	Pro	Ser	Glu
Leu 225	Asp	Αla	Leu	Thr	Pro 230	Leu	val	Thr	Ile	Ser 235	Gly	Thr	Asp	Thr	Ser 240
Thr	Thr	Phe	Pro	Thr 245	Leu	Thr	Lys	Ser	Pro 250	His	Glu	Thr	Glu	Thr 255	Arg
Thr	Thr	тгр	Leu 260	Thr	His	Pro	Ala	G]u 265	Thr	Ser	Ser	Thr	11e 270	Pro	Arg
Thr	Ile	Pro 275	Asn	Phe	Ser	His	ніs 280	Glu	Ser	Asp	Ala	Thr 285	Pro	Ser	Ile
Ala	Thr 290	Ser	Pro	Gly	Ala	G]u 295	Thr	Ser	Ser	Ala	Ile 300	Pro	Ile	Met	Thr
va1 305	Ser	Pro	Gly	Ala	Glu 310	Asp	Leu	Val	Thr	Ser 315	Gln	val	Thr	Ser	Ser 320
Gly	Thr	Asp	Arg	Asn 325	Met	Thr	Ile	Pro	Thr 330	Leu	Thr	Leu	Ser	Pro 335	Gly
Glu	Pro	Lys	Thr 340	Ile	Ala	Ser	Leu	va1 345	Thr	His	Pro	Glu	Ala 350	Gln	Thr
Ser	Ser	Ala 355	Ile	Pro	Thr	Ser	Thr 360	Ile	Ser	Pro	Ala	va1 365	Ser	Arg	Leu
val	Thr 370	Ser	Met	val	Thr	Ser 375	Leu	Ala	Ala	Lys	Thr 380	Ser	Thr	Thr	Asn
Arg 385	Ala	Leu	Thr	Asn	Ser 390	Pro	Gly	Glu	Pro	Ala 395	Thr	Thr	٧a٦	Ser	Leu 400

Val Thr His Pro Ala Gln Thr Ser Pro Thr Val Pro Trp Thr Thr Ser 405 Ile Phe Phe His Ser Lys Ser Asp Thr Thr Pro Ser Met Thr Thr Ser His Gly Ala Glu Ser Ser Ser Ala Val Pro Thr Pro Thr Val Ser Thr Glu Val Pro Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val 455 Ile Ser Thr Thr Ile Pro Ile Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Glu Glu Ala Ser Ser Ala Ile Pro Thr Pro Thr Val Ser Pro Gly Val Pro Gly Val Val Thr Ser 500 505 510 Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr Phe Ser Leu Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser 540 530 His Gly Thr Glu Ala Gly Ser Ala Val Pro Thr Val Leu Pro Glu Val 545 550 555 560 Pro Gly Met Val Thr Ser Leu Val Ala Ser Ser Arg Ala Val Thr Ser 565 570 575 Thr Thr Leu Pro Thr Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Thr Val Pro 600 Thr Val Ser Pro Glu Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Ser Gly Val Asn Ser Thr Ser Ile Pro Thr Leu Ile Leu Ser Pro Gly Glu Leu Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Page 182

Ala Ser Ser Ala Val Pro Thr Pro Thr Val Ser Pro Gly Val Ser Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr Leu Ser Ser Ser Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Val Glu Ala Ser Ser Ala Val Leu Thr Val 705 710 715 720 Ser Pro Glu Val Pro Gly Met Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Thr Leu Thr Ile Ser Ser Asp Glu 740 745 750 Pro Glu Thr Thr Thr Ser Leu Val Thr His Ser Glu Ala Lys Met Ile Ser Ala Ile Pro Thr Leu Ala Val Ser Pro Thr Val Gln Gly Leu Val 770 780 Thr Ser Leu Val Thr Ser Ser Gly Ser Glu Thr Ser Ala Phe Ser Asn 785 790 795 800 Leu Thr Val Ala Ser Ser Gln Pro Glu Thr Ile Asp Ser Trp Val Ala His Pro Gly Thr Glu Ala Ser Ser Val Val Pro Thr Leu Thr Val Ser 820 825 830 Thr Gly Glu Pro Phe Thr Asn Ile Ser Leu Val Thr His Pro Ala Glu Ser Ser Ser Thr Leu Pro Arg Thr Thr Ser Arg Phe Ser His Ser Glu 850 855 860 Leu Asp Thr Met Pro Ser Thr Val Thr Ser Pro Glu Ala Glu Ser Ser Ser Ala Ile Ser Thr Thr Ile Ser Pro Gly Ile Pro Gly Val Leu Thr 885 890 895 Ser Leu Val Thr Ser Ser Gly Arg Asp Ile Ser Ala Thr Phe Pro Thr Page 183

Val Pro Glu Ser Pro His Glu Ser Glu Ala Thr Ala Ser Trp Val Thr 915 920 925

His Pro Ala Val Thr Ser Thr Thr Val Pro Arg Thr Thr Pro Asn Tyr 930 935 940

Ser His Ser Glu Pro Asp Thr Thr Pro Ser Ile Ala Thr Ser Pro Gly 945 955 960

Ala Glu Ala Thr Ser Asp Phe Pro Thr Ile Thr Val Ser Pro Asp Val 965 970 975

Pro Asp Met Val Thr Ser Gln Val Thr Ser Ser Gly Thr Asp Thr Ser 980 985 990

Ile Thr Ile Pro Thr Leu Thr Leu Ser Ser Gly Glu Pro Glu Thr Thr 995 1000 1005

Thr Ser Phe Ile Thr Tyr Ser Glu Thr His Thr Ser Ser Ala Ile 1010 1015 1020

Pro Thr Leu Pro Val Ser Pro Gly Ala Ser Lys Met Leu Thr Ser 1025 1030 1035

Leu Val Ile Ser Ser Gly Thr Asp Ser Thr Thr Thr Phe Pro Thr 1040 1045 1050

Leu Thr Glu Thr Pro Tyr Glu Pro Glu Thr Thr Ala Ile Gln Leu 1055 1060 1065

Ile His Pro Ala Glu Thr Asn Thr Met Val Pro Arg Thr Thr Pro 1070 1080

Lys Phe Ser His Ser Lys Ser Asp Thr Thr Leu Pro Val Ala Ile 1085 1090 1095

Thr Ser Pro Gly Pro Glu Ala Ser Ser Ala Val Ser Thr Thr 1100 1105 1110

Ile Ser Pro Asp Met Ser Asp Leu Val Thr Ser Leu Val Pro Ser 1115 1120 1125

Ser Gly Thr Asp Thr Ser Thr Thr Phe Pro Thr Leu Ser Glu Thr 1130 1140

09-965738substitute.ST25.txt Pro Tyr Glu Pro Glu Thr Thr Ala Thr Trp Leu Thr His Pro Ala Glu Thr Ser Thr Thr Val Ser Gly Thr Ile Pro Asn Phe Ser His Gly Ser Asp Thr Ala Pro Ser Met Val Thr Ser Pro Gly Val 1175 1180 1185 Asp Thr Arg Ser Gly Val Pro Thr Thr Thr Ile Pro Pro Ser Ile Pro Gly Val Val Thr Ser Gln Val Thr Ser Ser Ala Thr Asp Thr 1205 1210 1215 Ser Thr Ala Ile Pro Thr Leu Thr Pro Ser Pro Gly Glu Pro Glu 1220 1225 Thr Thr Ala Ser Ser Ala Thr His Pro Gly Thr Gln Thr Gly Phe 1240 1235 Thr Val Pro Ile Arg Thr Val Pro Ser Ser Glu Pro Asp Thr Met Ala Ser Trp Val Thr His Pro Pro Gln Thr Ser Thr Pro Val Ser 1265 1270 1275 Arg Thr Thr Ser Ser Phe Ser His Ser Ser Pro Asp Ala Thr Pro Val Met Ala Thr Ser Pro Arg Thr Glu Ala Ser Ser Ala Val Leu 1305 Thr Thr Ile Ser Pro Gly Ala Pro Glu Met Val Thr Ser Gln Ile Thr Ser Ser Gly Ala Ala Thr Ser Thr Thr Val Pro Thr Leu Thr 1335 His Ser Pro Gly Met Pro Glu Thr Thr Ala Leu Leu Ser Thr His 1340 Pro Arg Thr Glu Thr Ser Lys Thr Phe Pro Ala Ser Thr Val Phe 1355 Pro Gln Val Ser Glu Thr Thr Ala Ser Leu Thr Ile Arg Pro Gly 1380 1370 Page 185

Ala Glu Thr Ser Thr Ala Leu Pro Thr Gln Thr Thr Ser Ser Leu 1385 1390 1395

Phe Thr Leu Leu Val Thr Gly Thr Ser Arg Val Asp Leu Ser Pro 1400 1410

Thr Ala Ser Pro Gly Val Ser Ala Lys Thr Ala Pro Leu Ser Thr 1415 1420 1425

His Pro Gly Thr Glu Thr Ser Thr Met Ile Pro Thr Ser Thr Leu 1430 1440

Ser Leu Gly Leu Leu Glu Thr Thr Gly Leu Leu Ala Thr Ser Ser 1445 1450 1455

Ser Ala Glu Thr Ser Thr Ser Thr Leu Thr Leu Thr Val Ser Pro 1460 1470

Ala Val Ser Gly Leu Ser Ser Ala Ser Ile Thr Thr Asp Lys Pro 1475 1480 1485

Gln Thr Val Thr Ser Trp Asn Thr Glu Thr Ser Pro Ser Val Thr 1490 1495 1500

Ser Val Gly Pro Pro Glu Phe Ser Arg Thr Val Thr Gly Thr Thr 1505 1510 1515

Met Thr Leu Ile Pro Ser Glu Met Pro Thr Pro Pro Lys Thr Ser 1520 1525 1530

His Gly Glu Gly Val Ser Pro Thr Thr Ile Leu Arg Thr Thr Met
1535 1540 1545

Val Glu Ala Thr Asn Leu Ala Thr Thr Gly Ser Ser Pro Thr Val 1550 1560

Ala Lys Thr Thr Thr Phe Asn Thr Leu Ala Gly Ser Leu Phe 1565 1570 1575

Thr Pro Leu Thr Thr Pro Gly Met Ser Thr Leu Ala Ser Glu Ser 1580 1585 1590

Val Thr Ser Arg Thr Ser Tyr Asn His Arg Ser Trp Ile Ser Thr 1595 1600 1605

Thr Ser Ser Tyr Asn Arg Arg Tyr Trp Thr Pro Ala Thr Ser Thr 1610 1615 1620 Page 186

Pro Val Thr Ser Thr Phe Ser Pro Gly Ile Ser Thr Ser Ser Ile Pro Ser Ser Thr Ala Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Glu Leu Gln 1680 1675 1670 Gly Leu Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu 1685 1695 Tyr Ser Gly Cys Arg Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro 1715 1720 1725 Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser 1730 1740 Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp 1745 1750 1755 Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly Thr 1775 1780 1785 Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr 150 <210> <211> 156 <212> PRT <213> Homo sapiens

<400> 150

Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe 20 25 30 Page 187

Asn	Thr	Thr 35	Glu	Arg	val	Leu	G]n 40	Gly	Leu	Leu	Lys	Pro 45	Leu	Phe	Lys	
Asn	Thr 50	Ser	val	Gly	Pro	Leu 55	Tyr	Ser	Gly	Cys	Arg 60	Leu	Thr	Leu	Leu	
Arg 65	Pro	Glu	Lys	His	G1u 70	Αla	Αla	Thr	GТу	va1 75	Asp	Thr	Ile	Cys	Thr 80	
His	Arg	٧al	Asp	Pro 85	Ile	Glу	Pro	Glу	Leu 90	Asp	Arg	Glu	Arg	Leu 95	Tyr	
тгр	Glu	Leu	Ser 100	Gln	Leu	Thr	Asn	Ser 105	Ile	Thr	Glu	Leu	Gly 110	Pro	Tyr	
Thr	Leu	Asp 115	Arg	Asp	Ser	Leu	Tyr 120	val	Asn	Gly	Phe	Asn 125	Pro	Arg	Ser	
Ser	val 130	Pro	Thr	Thr	Ser	Thr 135	Pro	Gly	Thr	Ser	Thr 140	val	His	Leu	Ala	
Thr 145	Ser	Glу	Thr	Pro	Ser 150	Ser	Leu	Pro	Lys	Leu 155	Thr					
<210 <211 <212 <213	L> : 2> [L51 507 DNA Homo	sapi	iens												
<220 <221 <222	L> (DS (1)	. (507	')												
<400 atg Met 1	aga	L51 gga Gly	tcg Ser	cat His 5	cac His	cat His	cac His	cat His	cac His 10	gga Gly	tcc Ser	atg Met	ggc Gly	cac His 15	aca Thr	48
gag Glu	cct Pro	ggc Gly	cct Pro 20	ctc Leu	ctg Leu	ata Ile	cca Pro	ttc Phe 25	act Thr	ttc Phe	aac Asn	ttt Phe	acc Thr 30	atc Ile	acc Thr	96
aac Asn	ctg Leu	cat His 35	tat Tyr	gag Glu	gaa Glu	aac Asn	atg Met 40	caa Gln	cac His	cct Pro	ggt Gly	tcc Ser 45	agg Arg	aag Lys	ttc Phe	144
aac Asn	acc Thr 50	acg Thr	gag Glu	agg Arg	gtt Val	ctg Leu 55	cag Gln	ggt Gly	ctg Leu	ctc Leu	aag Lys 60	ccc Pro	ttg Leu	ttc Phe	aag Lys	192
aac Asn	acc Thr	agt Ser	gtt val	ggc Gly	cct Pro	ctg Leu	tac Tyr	tct Ser	Gly	tgc Cys age :	Arg	ctg Leu	acc Thr	ttg Leu	ctc Leu	240

65					70		09-9	96573	88suk	ostii 75	ute.	ST25	5.txt	=	80	
aga Arg	cct Pro	gag Glu	aag Lys	cat His 85	gag Glu	gca Ala	gcc Ala	act Thr	gga Gly 90	gtg Val	gac Asp	acc Thr	atc Ile	tgt Cys 95	acc Thr	288
cac His	cgc Arg	gtt Val	gat Asp 100	ccc Pro	atc Ile	gga Gly	cct Pro	gga Gly 105	ctg Leu	gac Asp	aga Arg	gag Glu	cgg Arg 110	cta Leu	tac Tyr	336
tgg Trp	gag Glu	ctg Leu 115	agc Ser	cag Gln	ctg Leu	acc Thr	aac Asn 120	agc Ser	atc Ile	aca Thr	gag Glu	ctg Leu 125	gga Gly	ccc Pro	tac Tyr	384
acc Thr	ctg Leu 130	gac Asp	agg Arg	gac Asp	agt Ser	ctc Leu 135	tat Tyr	gtc val	aat Asn	ggc Gly	ttc Phe 140	aac Asn	cct Pro	cgg Arg	agc Ser	432
tct Ser 145	gtg Val	cca Pro	acc Thr	acc Thr	agc Ser 150	act Thr	cct Pro	ggg Gly	acc Thr	tcc Ser 155	aca Thr	gtg Val	cac His	ctg Leu	gca Ala 160	480
acc Thr	tct Ser	ggg Gly	act Thr	cca Pro 165	tcc Ser	tcc Ser	ctg Leu	cct Pro								507
<210 <211 <212 <213	L>	L52 L69 PRT Homo	sap [*]	iens												
<400)> :	L52														
_			Ser	His 5	His	His	His	His	ніs 10	Gly	Ser	Met	Gly	ніs 15	Thr	
Met 1	Arg	Gly		5	ніs Leu				10					15		
Met 1 Glu	Arg Pro	Gly Gly	Pro 20	5 Leu		Ile	Pro	Phe 25	10 Thr	Phe	Asn	Phe	Thr 30	15 Ile	Thr	
Met 1 Glu Asn	Arg Pro Leu	Gly Gly His 35	Pro 20 Tyr	5 Leu Glu	Leu	Ile Asn	Pro Met 40	Phe 25 Gln	Thr	Phe Pro	Asn Gly	Phe Ser 45	Thr 30 Arg	Ile Lys	Thr Phe	
Met 1 Glu Asn	Arg Pro Leu Thr	Gly Gly His 35 Thr	Pro 20 Tyr Glu	5 Leu Glu Arg	Leu Glu	Ile Asn Leu 55	Pro Met 40 Gln	Phe 25 Gln Gly	Thr His Leu	Phe Pro Leu	Asn Gly Lys 60	Phe Ser 45 Pro	Thr 30 Arg Leu	Ile Lys Phe	Thr Phe Lys	
Met 1 Glu Asn Asn	Arg Pro Leu Thr 50	Gly Gly His 35 Thr	Pro 20 Tyr Glu Val	5 Leu Glu Arg Gly	Leu Glu Val	Ile Asn Leu 55	Pro Met 40 Gln Tyr	Phe 25 Gln Gly Ser	Thr His Leu Gly	Phe Pro Leu Cys 75	Asn Gly Lys 60 Arg	Phe Ser 45 Pro Leu	Thr 30 Arg Leu Thr	Ile Lys Phe Leu	Thr Phe Lys Leu 80	
Met 1 Glu Asn Asn Asn 65	Arg Pro Leu Thr 50 Thr	Gly His 35 Thr Ser	Pro 20 Tyr Glu Val	5 Leu Glu Arg Gly His	Leu Glu Val Pro 70	Ile Asn Leu 55 Leu	Pro Met 40 Gln Tyr	Phe 25 Gln Gly Ser	Thr His Leu Gly Gly 90	Phe Pro Leu Cys 75	Asn Gly Lys 60 Arg	Phe Ser 45 Pro Leu Thr	Thr 30 Arg Leu Thr	Ile Lys Phe Leu Cys	Thr Phe Lys Leu 80	

Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser 130 140

Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala 145 150 155 160

Thr Ser Gly Thr Pro Ser Ser Leu Pro 165

<210> 153 <211> 507 <212> DNA

<213> Homo sapiens

<400> 153 aggcagggag gatggagtcc cagaggttgc caggtgcact gtggaggtcc caggagtgct 60 120 ggtggttggc acagagctcc gagggttgaa gccattgaca tagagactgt ccctgtccag 180 qqtqtaqqqt cccaqctctg tgatgctgtt ggtcagctgg ctcagctccc agtatagccg 240 ctctctqtcc agtccaggtc cgatgggatc aacgcggtgg gtacagatgg tgtccactcc 300 agtggctgcc tcatgcttct caggtctgag caaggtcagt ctgcagccag agtacagagg 360 qccaacactg gtgttcttga acaagggctt gagcagaccc tgcagaaccc tctccgtggt 420 gttgaacttc ctggaaccag ggtgttgcat gttttcctca taatgcaggt tggtgatggt 480 aaaqttqaaa gtgaatggta tcaggagagg gccaggctct gtgtggccca tggatccgtg 507 atggtgatgg tgatgcgatc ctctcat

<210> 154 <211> 9 <212> PRT <213> Homo sapiens

Are Low Two Trop Clu Low Son Cln I

Arg Leu Tyr Trp Glu Leu Ser Gln Leu 1 5

<210> 155 <211> 9 <212> PRT <213> Homo sap

<213> Homo sapiens

<400> 155

<400> 154

Thr Leu Asp Arg Asp Ser Leu Tyr Val

<210> 156

<211>

<212> PRT

<213> Homo sapiens

<400> 156

Val Leu Gln Gly Leu Leu Lys Pro Leu 1 5

<210> 157

<211> 9

<212> PRT

<213> Homo sapiens

<400> 157

Gln Leu Thr Asn Ser Ile Thr Glu Leu 5

<210> 158

780

PRT

<211> <212> <213> Homo sapiens

<400> 158

Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Ala Thr Glu Arg Glu Leu Gln Gly Leu Leu Lys Pro Leu Phe Arg 35 40 45

Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly Cys Arg Leu Ala Ser Leu 50 60

Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr 65 70 75 80

His Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr 85 90 95

Trp Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr 100 105 110

Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser

Ser Met Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Val Gly 130

Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr Ala Ala Gly Pro Leu Leu Met Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu 180 185 190 Ser Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val 195 200 205 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 210 220 Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp 225 230 235 240 Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser 245 250 255 Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg 260 265 270 Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr 275 280 285 Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr 290 295 300 Pro Ser Ser Leu Ser Ser Pro Thr Ile Met Ala Gly Pro Leu Leu Val 305 310 315 320 305 Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp 325 330 335 Met Gly His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 340 345 350 Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser Val Gly Pro Leu 355 360 365 Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys Asp Gly Ala 370 380 Ala Thr Gly Val Asp Ala Ile Cys Ile His His Leu Asp Pro Lys Ser 390 395 Page 192

Pro Gly Leu Asn Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr 405 410 415 Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 420 425 430 Tyr Val Asn Gly Phe Thr His Arg Thr Ser Val Pro Thr Ser Ser Thr 435 440 445 Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe Ser 450 460 Leu Pro Ser Pro Ala Thr Ala Gly Pro Leu Leu Val Leu Phe Thr Leu 465 470 475 480 Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp Met His Arg Pro 485 490 495 Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Thr Leu Leu 500 510Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys 515 520 525 Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val 530 540 Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp 545 550 555 560 Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys 565 570 575 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 580 585 590 Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Ser 595 600 605 Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr 610 620 Ala Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 630 Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe Page 193

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys 660 665 670 660

Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu 675 680 685

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr 690 695 700

His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr 705 710 715 720

Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr 725 730 735

Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr 740 745 750

Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly 755 760 765

Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr 770 775 780

<210> 159 <211> 780

<212> PRT Homo sapiens <213>

<400> 159

Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys 35 40 45

Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu 50 60

Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser 65 70 75 80

His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Page 194

Trp Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr 100 105 110 100 Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser 115 120 125 Ser Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 180 185 190 Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val 195 200 205 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys 210 220 Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn 225 230 235 240 Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser 245 250 255 Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg 260 265 270 Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr 275 280 285 Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr 290 295 300 Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro 305 310 315 320 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met 325 330 335

09-965738substitute.ST25.txt His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu Gln 340 345 350 Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr 355 360 365 Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala 370 375 380 Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg Pro 385 390 400 Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro 435 440 445 Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe 450 460 Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn 465 470 475 480 Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly 485 490 495 Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys 500 505 Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg 515 520 525 Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp 530 535 540 Ala Val Cys Leu Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg 545 550 555 560 Glu Gln Leu Tyr Cys Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu 565 570 575 Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe 580 585

Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr 595 600 605

Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr 610 620

Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr 625 630 635 640

Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe 645 650 655

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys 660 665 670

Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu 675 680 685

Arg Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr 690 695 700

His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr 705 710 715 720

Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr
725 730 735

Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser 740 745 750

Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala 755 760 765

Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr 770 775 780

<400> 160

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Arg Phe 20 25 30

<210> 160

<211> 624

<212> PRT

<213> Homo sapiens

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Lys
35 40 45 Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu 50 60 Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr 65 70 75 80 His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr 85 90 95 Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr 100 105 110 Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser 115 120 125 Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala 130 135 140 Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asp Leu His Tyr 165 170 175 Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 180 185 190 Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val 195 200 205 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 210 220 His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp 225 230 235 240 Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser 245 250 255 Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg 260 265 270 Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr 275 280 285 Page 198

Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr 290 295 300 Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val 305 310 315 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met 325 330 335 Arg His Pro Gly Ser Arg Lys Phe Ser Thr Thr Glu Arg Val Leu Gln 340 345 350 Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala 370 375 380 Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser Pro 385 390 395 Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr Arg Thr Pro 435 440 445 Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu 450 460 Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn 465 470 Phe Thr Ile Thr Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly 495 Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg 500 505 510 Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg 515 520 525 Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Page 199

Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg 545 550 555 560

Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu 565 570 575

Leu Gly Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe 580 585

Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala 595 600 605

Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr 610 615 620

<210> 161

<211> 468

<212> PRT

<213> Homo sapiens

<400> 161

Ala Thr Gly Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Met Pro Leu Phe Lys 35 40 45

Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu 50 60

Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys Thr 65 70 75 80

His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr 85 90 95

Trp Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr 100 105 110

Thr Leu Asp Arg His Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser 115 120 125

Ser Met Thr Thr Arg Thr Pro Asp Thr Ser Thr Met His Leu Ala Page 200

Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Leu Arg Tyr 165 170 175 Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 180 185 190 Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val 195 200 205 Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys 210 220 Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp 225 230 235 240 Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser 245 250 255 Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Gln Asp Arg 260 265 270Asp Ser Leu Tyr Asn Val Gly Phe Thr Gln Arg Ser Ser Val Pro Thr 275 280 285 Thr Ser Val Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr 290 295 300 Pro Val Ser Lys Pro Gly Pro Ser Ala Ala Ser Pro Leu Leu Val Leu 305 310 315 320 Phe Thr Leu Asn Gly Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met 325 330 335 Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln 340 345 350 Gly Leu Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr 355 360 365 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Thr Ala 370 380

```
09-965738substitute.ST25.txt
Thr Gly Val Asp Ala Ile Cys Thr His His Pro Asp Pro Lys Ser Pro
                      390
                                             395
Arg Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His
Asn Ile Thr Glu Leu Gly His Tyr Ala Leu Asp Asn Asp Ser Leu Phe
Val Asn Gly Phe Thr His Arg Ser Ser Val Ser Thr Thr Ser Thr Pro
435 440 445
Gly Thr Pro Thr Val Tyr Leu Gly Ala Ser Lys Thr Pro Ala Ser Ile
Phe Gly Pro Ser
<210>
        162
        11721
<211>
<212>
        PRT
<213>
       Homo sapiens
<220>
<221>
       MISC_FEATURE
       (1)..(11721)
Any "X" = any amino acid
<222>
<223>
<400>
Met Glu His Ile Thr Lys Ile Pro Asn Glu Ala Ala His Arg Gly Thr 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Ile Arg Pro Val Lys Gly Pro Gln Thr Ser Thr Ser Pro Ala Ser Pro
Lys Gly Leu His Thr Gly Gly Thr Lys Arg Met Glu Thr Thr Thr \frac{1}{35} 40 45
Ala Leu Lys Thr Thr Thr Ala Leu Lys Thr Thr Ser Arg Ala Thr
Leu Thr Thr Ser Val Tyr Thr Pro Thr Leu Gly Thr Leu Thr Pro Leu 65 70 75 80
Asn Ala Ser Arg Gln Met Ala Ser Thr Ile Leu Thr Glu Met Met Ile
Thr Thr Pro Tyr Val Phe Pro Asp Val Pro Glu Thr Thr Ser Ser Leu
                                    105
```

Page 202

Ala Thr Ser Leu Gly Ala Glu Thr Ser Thr Ala Leu Pro Arg Thr Thr Ser Val Leu Asn Arg Glu Ser Glu Thr Thr Ala Ser Leu Val Ser Arg Ser Gly Ala Glu Arg Ser Pro Val Ile Gln Thr Leu Asp Val Ser Ser Ser Glu Pro Asp Thr Thr Ala Ser Trp Val Ile His Pro Ala Glu 165 170 175 Thr Ile Pro Thr Val Ser Lys Thr Thr Pro Asn Phe Phe His Ser Glu Leu Asp Thr Val Ser Ser Thr Ala Thr Ser His Gly Ala Asp Val Ser Ser Ala Ile Pro Thr Asn Ile Ser Pro Ser Glu Leu Asp Ala Leu Thr Pro Leu Val Thr Ile Ser Gly Thr Asp Thr Ser Thr Thr Phe Pro Thr Leu Thr Lys Ser Pro His Glu Thr Glu Thr Arg Thr Thr Trp Leu Thr His Pro Ala Glu Thr Ser Ser Thr Ile Pro Arg Thr Ile Pro Asn Phe 260 265 270 Ser His His Glu Ser Asp Ala Thr Pro Ser Ile Ala Thr Ser Pro Gly 285 Ala Glu Thr Ser Ser Ala Ile Pro Ile Met Thr Val Ser Pro Gly Ala Glu Asp Leu Val Thr Ser Gln Val Thr Ser Ser Gly Thr Asp Arg Asn 310 Met Thr Ile Pro Thr Leu Thr Leu Ser Pro Gly Glu Pro Lys Thr Ile Ala Ser Leu Val Thr His Pro Glu Ala Gln Thr Ser Ser Ala Ile Pro 340 Thr Ser Thr Ile Ser Pro Ala Val Ser Arg Leu Val Thr Ser Met Val Page 203

Thr Ser Leu Ala Ala Lys Thr Ser Thr Thr Asn Arg Ala Leu Thr Asn 370 380 Ser Pro Gly Glu Pro Ala Thr Thr Val Ser Leu Val Thr His Pro Ala Gln Thr Ser Pro Thr Val Pro Trp Thr Thr Ser Ile Phe Phe His Ser Lys Ser Asp Thr Thr Pro Ser Met Thr Thr Ser His Gly Ala Glu Ser 420 425 430 Ser Ser Ala Val Pro Thr Pro Thr Val Ser Thr Glu Val Pro Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Ile Ser Thr Thr Ile Pro Ile Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Glu Glu Ala Ser Ser Ala Ile Pro Thr Pro Thr 485 490 495 Val Ser Pro Gly Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser 500 510 Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr Phe Ser Leu Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Thr Glu Ala Gly Ser Ala Val Pro Thr Val Leu Pro Glu Val Pro Gly Met Val Thr Ser Leu Val Ala Ser Ser Arg Ala Val Thr Ser Thr Thr Leu Pro Thr 565 570 575 Leu Thr Leu Ser Pro Gly Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Thr Val Pro Thr Val Ser Pro Glu 595 600 . 605

09-965738substitute.ST25.txt Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Ser Gly Val Asn Ser Thr Ser Ile Pro Thr Leu Ile Leu Ser Pro Gly Glu Leu Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Ala Val Pro Thr Pro Thr Val Ser Pro Gly Val Ser Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr 680 685 Leu Ser Ser Ser Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His 690 Gly Val Glu Ala Ser Ser Ala Val Leu Thr Val Ser Pro Glu Val Pro 705 710 715 720 Gly Met Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr 725 730 735 Thr Ile Pro Thr Leu Thr Ile Ser Ser Asp Glu Pro Glu Thr Thr Thr 740 745 750 Ser Leu Val Thr His Ser Glu Ala Lys Met Ile Ser Ala Ile Pro Thr Leu Ala Val Ser Pro Thr Val Gln Gly Leu Val Thr Ser Leu Val Thr Ser Ser Gly Ser Glu Thr Ser Ala Phe Ser Asn Leu Thr Val Ala Ser Ser Gln Pro Glu Thr Ile Asp Ser Trp Val Ala His Pro Gly Thr Glu Ala Ser Ser Val Val Pro Thr Leu Thr Val Ser Thr Gly Glu Pro Phe 820 830 Thr Asn Ile Ser Leu Val Thr His Pro Ala Glu Ser Ser Ser Thr Leu 845 835 Pro Arg Thr Thr Ser Arg Phe Ser His Ser Glu Leu Asp Thr Met Pro Page 205

- Ser Thr Val Thr Ser Pro Glu Ala Glu Ser Ser Ser Ala Ile Ser Thr 865 870 875 880
- Thr Ile Ser Pro Gly Ile Pro Gly Val Leu Thr Ser Leu Val Thr Ser 885 890 895
- Ser Gly Arg Asp Ile Ser Ala Thr Phe Pro Thr Val Pro Glu Ser Pro 900 905 910
- His Glu Ser Glu Ala Thr Ala Ser Trp Val Thr His Pro Ala Val Thr 915 920 925
- Ser Thr Thr Val Pro Arg Thr Thr Pro Asn Tyr Ser His Ser Glu Pro 930 935 940
- Asp Thr Thr Pro Ser Ile Ala Thr Ser Pro Gly Ala Glu Ala Thr Ser 945 950 955 960
- Asp Phe Pro Thr Ile Thr Val Ser Pro Asp Val Pro Asp Met Val Thr 965 970 975
- Ser Gln Val Thr Ser Ser Gly Thr Asp Thr Ser Ile Thr Ile Pro Thr 980 985 990
- Leu Thr Leu Ser Ser Gly Glu Pro Glu Thr Thr Ser Phe Ile Thr 995 1000 1005
- Tyr Ser Glu Thr His Thr Ser Ser Ala Ile Pro Thr Leu Pro Val 1010 1015 1020
- Ser Pro Gly Ala Ser Lys Met Leu Thr Ser Leu Val Ile Ser Ser 1025 1030 1035
- Gly Thr Asp Ser Thr Thr Thr Phe Pro Thr Leu Thr Glu Thr Pro 1040 1045 1050
- Tyr Glu Pro Glu Thr Thr Ala Ile Gln Leu Ile His Pro Ala Glu 1055 1060 1065
- Thr Asn Thr Met Val Pro Arg Thr Thr Pro Lys Phe Ser His Ser 1070 1075 1080
- Lys Ser Asp Thr Thr Leu Pro Val Ala Ile Thr Ser Pro Gly Pro 1085 1090 1095
- Glu Ala Ser Ser Ala Val Ser Thr Thr Ile Ser Pro Asp Met 1100 1105 1110 Page 206

Page 207

Ser Asp_ Leu Val Thr Ser Leu Val Pro Ser Ser Gly_ Thr Asp Thr 1115 Ser Thr Thr Phe Pro Thr Leu Ser Glu Thr Pro Tyr Glu Pro Glu Thr Thr Ala Thr Trp Leu Thr His Pro Ala Glu Thr Ser Thr Thr Val Ser Gly Thr Ile Pro Asn Phe Ser His Arg Gly Ser Asp Thr 1160 1165 11701160 Ala Pro Ser Met Val Thr Ser Pro Gly Val Asp Thr Arg Ser Gly 1175 1180 Val Pro Thr Thr Thr Ile Pro Pro Ser Ile Pro Gly Val Val Thr 1190 1195 Ser Gln Val Thr Ser Ser Ala Thr Asp Thr Ser Thr Ala Ile Pro 1205 Thr Leu Thr Pro Ser Pro Gly Glu Pro Glu Thr Thr Ala Ser Ser 1220 1230 Ala Thr His Pro Gly Thr Gln Thr Gly Phe Thr Val Pro Ile Arg 1235 1240 1245 Thr Val Pro Ser Ser Glu Pro Asp Thr Met Ala Ser Trp Val Thr His Pro Pro Gln Thr Ser Thr Pro Val Ser Arg Thr Thr Ser Ser 1275 1270 Phe Ser His Ser Ser Pro Asp Ala Thr Pro Val Met Ala Thr Ser Pro Arg Thr Glu Ala Ser Ser Ala Val Leu Thr Thr Ile Ser Pro 1305 1300 Gly Ala Pro Glu Met Val Thr Ser Gln Ile Thr Ser Ser Gly Ala 1320 1310 Ala Thr Ser Thr Thr Val Pro Thr Leu Thr His Ser Pro Gly Met 1335 1325 Pro Glu Thr Thr Ala Leu Leu Ser Thr His Pro Arg Thr Glu Thr

1340 1345 1350 Ser Lys Thr Phe Pro Ala Ser Thr Val Phe Pro Gln Val Ser Glu Thr Thr Ala Ser Leu Thr Ile Arg Pro Gly Ala Glu Thr Ser Thr 1370 1380 Ala Leu Pro Thr Gln Thr Thr Ser Ser Leu Phe Thr Leu Leu Val 1390 Thr Gly Thr Ser Arg Val Asp Leu Ser Pro Thr Ala Ser Pro Gly 1400 1410 val Ser Ala Lys Thr Ala Pro Leu Ser Thr His Pro Gly Thr Glu Thr Ser Thr Met Ile Pro Thr Ser Thr Leu Ser Leu Gly Leu Leu Glu Thr Thr Gly Leu Leu Ala Thr Ser Ser Ser Ala Glu Thr Ser Thr Ser Thr Leu Thr Leu Thr Val Ser Pro Ala Val Ser Gly Leu 1470 1460 Ser Ser Ala Ser Ile Thr Thr Asp Lys Pro Gln Thr Val Thr Ser 1475 1480 Trp Asn Thr Glu Thr Ser Pro Ser Val Thr Ser Val Gly Pro Pro 1495 1500 1490 Glu Phe Ser Arg Thr Val Thr Gly Thr Thr Met Thr Leu Ile Pro

1505

Ser Glu Met Pro Thr Pro Pro Lys Thr Ser His Gly Glu Gly Val

Ser Pro Thr Thr Ile Leu Arg Thr Thr Met Val Glu Ala Thr Asn 1535 1540 1545 1535

Leu Ala Thr Thr Gly Ser Ser Pro Thr Val Ala Lys Thr Thr Thr

Thr Phe Asn Thr Leu Ala Gly Ser Leu Phe Thr Pro Leu Thr Thr 1565 1570 1575

											:e.ST2			
Pro	Gly 1580	Met	Ser	Thr	Leu	Ala 1585	Ser	Glu	Ser	Val	Thr 1590	Ser	Arg	Thr
Ser	Tyr 1595	Asn	His	Arg	Ser	Trp 1600	Ile	Ser	Thr	Thr	ser 1605	Ser	Tyr	Asn
Arg	Arg 1610	туг	Trp	Thr	Pro	Ala 1615	Thr	Ser	Thr	Pro	Val 1620	Thr	Ser	Thr
Phe	Ser 1625	Pro	Gly	Ile	Ser	Thr 1630	Ser	Ser	Ile	Pro	ser 1635	Ser	Thr	Ala
Ala	Thr 1640	val	Pro	Phe	Met	Val 1645	Pro	Phe	Thr	Leu	Asn 1650	Phe	Thr	Ile
Thr	Asn 1655	Leu	Gln	Tyr	Glu	Glu 1660	Asp	Met	Arg	His	Pro 1665	Gly	Ser	Arg
Lys	Phe 1670	Asn	Ala	Thr	Glu	Arg 1675	Glu	Leu	Gln	Gly	Leu 1680	Leu	Lys	Pro
Leu	Phe 1685	Arg	Asn	Ser	Ser	Leu 1690	Glu	Tyr	Leu	Tyr	Ser 1695	Gly	Cys	Arg
Leu	Ala 1700	Ser	Leu	Arg	Pro	Glu 1705	Lys	Asp	Ser	Ser	Ala 1710	Met	Ala	val
Asp	Ala 1715	Ile	Cys	Thr	His	Arg 1720	Pro	Asp	Pro	Glu	Asp 1725	Leu	Gly	Leu
Asp	Arg 1730	Glu	Arg	Leu	Tyr	Trp 1735	Glu	Leu	Ser	Asn	Leu 1740	Thr	Asn	GТу
Ile	Gln 1745	Glu	Leu	Gly	Pro	Tyr 1750	Thr	Leu	Asp	Arg	Asn 1755	Ser	Leu	Tyr
٧a٦	Asn 1760	Gly	Phe	Thr	His	Arg 1765	Ser	Ser	Met	Pro	Thr 1770	Thr	Ser	Thr
Pro	Gly 1775	Thr	Ser	Thr	۷a٦	Asp 1780	٧a٦	Gly	Thr	Ser	Gly 1785	Thr	Pro	Ser
Ser	Ser 1790	Pro	Ser	Pro	Thr	Ala 1795	Ala	Gly	Pro	Leu	Leu 1800	Met	Pro	Phe
Thr	Leu 1805	Asn	Phe	Thr	Ile	Thr 1810	Asn	Leu	Gln	Tyr	Glu 1815	Glu	Asp	Met
									Page	e 20	9			

Arg Arg Thr Gly Ser Arg Lys Phe Asn Thr Met Glu Ser Val Leu 1820 1830 Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro 1840 Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp 1850 1860 Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp 1865 1870 1875 Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu 1880 1885 1890 Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser 1915 val Ser Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg 1930 Thr Ser Gly Thr Pro Ser Ser Leu Ser Ser Pro Thr Ile Met Ala 1945 1950 1940 Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1955 Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys 1970 1980 1970 Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Ile Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu 2000 2010 Thr Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp 2015 2020 2025 Ala Ile Cys Ile His His Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile 2055 2050 Page 210

Lys	G1u 2060	Leu	Gly	Pro	Tyr	Thr 2065	Leu	Asp	Arg	Asn	ser 2070	Leu	Tyr	val
Asn	G1y 2075	Phe	Thr	His	Arg	Thr 2080	Ser	val	Pro	Thr	ser 2085	Ser	Thr	Pro
Glу	Thr 2090	Ser	Thr	val	Asp	Leu 2095	Gly	Thr	Ser	Gly	Thr 2100	Pro	Phe	Ser
Leu	Pro 2105	Ser	Pro	Ala	Thr	Ala 2110	Gly	Pro	Leu	Leu	val 2115	Leu	Phe	Thr
Leu	Asn 2120	Phe	Thr	Ile	Thr	Asn 2125	Leu	Lys	Tyr	Glu	Glu 2130	Asp	Met	His
Arg	Pro 2135	Gly	Ser	Arg	Lys	Phe 2140	Asn	Thr	Thr	Glu	Arg 2145	val	Leu	Gln
Thr	Leu 2150	Leu	Gly	Pro	Met	Phe 2155	Lys	Asn	Thr	Ser	val 2160	Gly	Leu	Leu
Tyr	Ser 2165	Gly	Cys	Arg	Leu	Thr 2170	Leu	Leu	Arg	Ser	Glu 2175	Lys	Asp	Gly
Ala	Ala 2180	Thr	Gly	٧a٦	Asp	Ala 2185	Ile	Cys	Thr	His	Arg 2190	Leu	Asp	Pro
Lys	Ser 2195	Pro	Gly	Leu	Asp	Arg 2200	Glu	Gln	Leu	Tyr	Trp 2205	Glu	Leu	Ser
Gln	Leu 2210	Thr	Asn	Gly	Ile	Lys 2215	Glu	Leu	Gly	Pro	Tyr 2220	Thr	Leu	Asp
Arg	Asn 2225	Ser	Leu	туг	val	Asn 2230	Gly	Phe	Thr	His	Trp 2235	Ile	Pro	val
Pro	Thr 2240	Ser	Ser	Thr	Pro	Gly 2245	Thr	Ser	Thr	Val	Asp 2250	Leu	Gly	Ser
Gly	Thr 2255	Pro	Ser	Ser	Leu	Pro 2260	Ser	Pro	Thr	Ala	Ala 2265	Gly	Pro	Leu
Leu	va1 2270	Pro	Phe	Thr	Leu	Asn 2275	Phe	Thr	Ile	Thr	Asn 2280	Leu	Gln	Tyr
Glu	Glu	Asp	Met	His	His	Pro	Gly	Ser	Arg Pag	Lys e 21	Phe 1	Asn	Thr	Thr

Glu Arg Val 2300	Leu Gln Gly	Leu Leu 2305	Gly Pro	Met Phe 2310	Lys Asn Thr
Ser Val Gly 2315	Leu Leu Tyr	Ser Gly 2320	Cys Arg	Leu Thr 2325	Leu Leu Arg
Ser Glu Lys 2330	Asp Gly Ala	Ala Thr 2335	Gly Val	Asp Ala 2340	Ile Cys Thr
His Arg Leu 2345	Asp Pro Lys	Ser Pro 2350	Gly Val	Asp Arg 2355	Glu Gln Leu
Tyr Trp Glu 2360	Leu Ser Gln	Leu Thr 2365	Asn Gly	Ile Lys 2370	Glu Leu Gly
Pro Tyr Thr 2375	Leu Asp Arg	Asn Ser 2380	Leu Tyr	Val Asn 2385	Gly Phe Thr
His Gln Thr 2390	Ser Ala Pro	Asn Thr 2395	Ser Thr	Pro Gly 2400	Thr Ser Thr
Val Asp Leu 2405	Gly Thr Ser	Gly Thr 2410	Pro Ser	Ser Leu 2415	Pro Ser Pro
Thr Ser Ala 2420	Gly Pro Leu	Leu Val 2425	Pro Phe	Thr Leu 2430	Asn Phe Thr
Ile Thr Asn 2435	Leu Gln Tyr	Glu Glu 2440	Asp Met	Arg His 2445	Pro Gly Ser
Arg Lys Phe 2450	Asn Thr Thr	Glu Arg 2455	Val Leu	Gln Gly 2460	Leu Leu Lys
Pro Leu Phe 2465	Lys Ser Thr	Ser Val 2470	Gly Pro	Leu Tyr 2475	Ser Gly Cys
Arg Leu Thr 2480	Leu Leu Arg	Ser Glu 2485	Lys Asp	Gly Ala 2490	Ala Thr Gly
Val Asp Ala 2495	Ile Cys Thr	His Arg 2500	Leu Asp	Pro Lys 2505	Ser Pro Gly
val Asp Arg 2510	Glu Gln Leu	Туг Тгр 2515	Glu Leu	Ser Gln 2520	Leu Thr Asn

09-965738substitute.ST25.txt Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu 2525 2530 2535 Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly Pro Leu Leu Val Pro 2570 2580 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 2585 2590 Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 2600 2605 2610 2600 Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly 2620 2615 Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 2630 2640 Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu 2645 2650 2655 Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu 2715 Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala 2720 Val Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn 2745 2735 Leu Gln Tyr Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Leu Phe 2765 2770 2775 2765 Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Ile 2780 2785 2790 Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala 2795 2800 2805 Ile Cys Thr His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg 2810 2815 2820 Glu Gln Leu Tyr Trp Gln Leu Ser Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val 2870 2875 2880 Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu 2885 2895 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His Arg 2900 2905 Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu Gln Gly 2915 2920 2925 Leu Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr 2930 2935 2940 2930 Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala 2945 2950 2955 Ala Thr Gly Met Asp Ala Val Cys Leu Tyr His Pro Asn Pro Lys 2970 Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln 2980 2985 Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg 3000 2995 Page 214

Asp	Ser 3005	Leu	Tyr	٧a٦	Asn	Gly 3010	Phe	Thr	ніѕ	Gln	Asn 3015	Ser	val	Pro
Thr	Thr 3020	Ser	Thr	Pro	Gly	Thr 3025	Ser	Thr	val	Tyr	Trp 3030	Ala	Thr	Thr
Gly	Thr 3035	Pro	Ser	Ser	Phe	Pro 3040	Gly	His	Thr	Glu	Pro 3045	Gly	Pro	Leu
Leu	Ile 3050	Pro	Phe	Thr	Phe	Asn 3055	Phe	Thr	Ile	Thr	Asn 3060	Leu	His	Tyr
Glu	G]u 3065	Asn	Met	Gln	His	Pro 3070		Ser	Arg	Lys	Phe 3075	Asn	Thr	Thr
Glu	Arg 3080	val	Leu	Gln	Gly	Leu 3085	Leu	Lys	Pro	Leu	Phe 3090	Lys	Asn	Thr
Ser	va1 3095	Gly	Pro	Leu	Tyr	Ser 3100	Gly	Cys	Arg	Leu	Thr 3105	Ser	Leu	Arg
Pro	Glu 3110	Lys	Asp	Gly	Ala	Ala 3115	Thr	Gly	Met	Asp	Ala 3120	val	Cys	Leu
Tyr	ніs 3125	Pro	Asn	Pro	Lys	Arg 3130	Pro	Gly	Leu	Asp	Arg 3135	Glu	Gln	Leu
Tyr	Cys 3140	Glu	Leu	Ser	Gln	Leu 3145	Thr	His	Asn	Ile	Thr 3150	Glu	Leu	Gly
Pro	Tyr 3155	Ser	Leu	Asp	Arg	Asp 3160	Ser	Leu	Tyr	val	Asn 3165	Gly	Phe	Thr
His	Gln 3170	Asn	Ser	val	Pro	Thr 3175	Thr	Ser	Thr	Pro	Gly 3180	Thr	Ser	Thr
∨al	Tyr 3185	Trp	Ala	Thr	Thr	Gly 3190	Thr	Pro	Ser	Ser	Phe 3195	Pro	Glу	His
Thr	G]u 3200	Pro	Gly	Pro	Leu	Leu 3205	Ile	Pro	Phe	Thr	Phe 3210	Asn	Phe	Thr
Ile	Thr 3215	Asn	Leu	His	Tyr	Glu 3220	Glu	Asn	Met	Gln	His 3225	Pro	Glу	Ser
Arg	Lys	Phe	Asn	Thr	Thr	Glu	Arg	val	Leu Page	Gln e 21	Gly 5	Leu	Leu	Lys

Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 3245 3250 3245 3250 Arg Leu Thr Leu Leu Arg Pro Glu Lys His Glu Ala Ala Thr Gly 3260 Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly 3275 3280 3285 Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn 3290 3295 3300 Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu 3305 3310 3315 Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro 3350 3360 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn 3370 3365 Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 3380 3390 Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly 3395 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 3410 3420 3410 His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val 3435 3425 3430 Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa

3460

09-965738substitute.ST25.txt Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa 3470 3475 3480 Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn 3525 3515 3520 Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe 3540 3530 3535 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe 3545 3550 3555 3550 3545 Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr 3560 3570 Leu Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala 3575 3580 3585 Ile Cys Ser His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg 3590 3595 3600 Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Ile Lys 3605 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn 3630 Gly Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu 3650 3660 Pro Ser Pro Thr Thr Ala Val Pro Leu Leu Val Pro Phe Thr Leu 3675 3665 3670 Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp Met Arg His 3680 3685 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 3695

Page 218

Leu Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr 3710 3715 3720 Ser Gly Cys Arg Leu Ile Ser Leu Arg Ser Glu Lys Asp Gly Ala 3725 3730 3735 Ala Thr Gly Val Asp Ala Ile Cys Thr His His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Gln Leu Ser Gln 3755 3760 3765 Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser_Leu Tyr Val Asn Gly_Phe Thr His Arg Ser_Ser Gly Leu Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu 3820 Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr 3840 3830 3835 Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr 3855 3845 Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Ser 3865 3870 3860 Ser val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu 3890 3895 3900 Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu 3905 3910 3915 3910 Tyr Trp Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr 3945 3940

His Gln Ser Ser Met Thr Thr Arg Thr Pro Asp Thr Ser Thr 3955 Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Cys Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser 40Ŏ5 4000 3995 Arg Lys Phe Asn Thr Met Glu Ser Val Leu Gln Gly Leu Leu Lys 4020 4010 4015 Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 4025 4035 4030 4025 Arg Leu Thr Leu Leu Arg Pro Lys Lys Asp Gly Ala Ala Thr Gly 4040 Val Asp Ala Ile Cys Thr His Arg Leu Asp Pro Lys Ser Pro Gly 4055 4060 4065 Leu Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Asn 4070 4080 Asp Ile Glu Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu **40**95 Tyr Val Asn Gly Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro 4120 Ser Ser Leu Ser Ser Pro Thr Ile Met Xaa Xaa Xaa Pro Leu Leu 4140 4130 4135 Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu 4150 4155 4145 Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 4160 4165 4170 Arg Val Leu Gln Gly Leu Leu Arg Pro Leu Phe Lys Asn Thr Ser Page 219

Val Ser Ser Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro 4190 4195 4200 Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Ala Cys Thr Tyr 4205 4210 4215 Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro 4240 Tyr Thr Leu Asp Arg Val Ser Leu Tyr Val Asn Gly Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile 4305 4295 4300 Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg 4310 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro 4325 Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly Cys Arg 4340 Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp Leu Gly Leu 4370 4380 Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr Asn Gly 4390 Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr

4405

09-965738substitute.ST25.txt Val Asn Gly Phe Thr His Arg Ser Ser Phe Leu Thr Thr Ser Thr 44Ž0 Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met 4460 4470 His Arg Pro Gly Ser Arg Arg Phe Asn Thr Thr Glu Arg Val Leu 4475 4480 4485 Gln Gly Leu Leu Thr Pro Leu Phe Lys Asn Thr Ser Val Gly Pro 4495 Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln 4510 Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val Asp 4520 4530 Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu 4535 4540 4545 Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val 4605 4595 4600 Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asp Leu 4610 4620 His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg_ Lys Phe Asn 4625 Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys 4640

Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu 4655 Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile 4670 4680 Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu 4690 Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu 4700 4710 Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro 4775 4780 4785 Gly Ser Arg Lys Phe Ser Thr Thr Glu Arg Val Leu Gln Gly Leu 4790 Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser 4810 4805 Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys Ser 4835 4840 4845 Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Lys Leu Ser Gln Leu 4850 4860 Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg His 4875 4870 Ser Leu Tyr Val Asn Gly Phe Thr His Gln Ser Ser Met Thr Thr 4890 4885 Page 222

Thr Arg_ Thr Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg 48**9**5 Thr Pro Ala Ser Leu Ser Gly Pro Thr Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 4940 4945 4950 Arg Val Leu Gln Gly Leu Leu Arg Pro Val Phe Lys_ Asn Thr Ser 4965 4955 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro 4970 4980 Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro 5020 5015 Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His 5035 Arg Ser Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val 5050 His Leu Glu Thr Ser Gly Thr Pro Ala Ser Leu Pro Gly His Thr 5065 5060 Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile 5075 5080 5085 Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg 5100 5095 5090 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro 5105 5115 Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Page 223

Leu Thr Leu Leu Arg Pro Glu Lys Arg Gly Ala Ala Thr Gly Val 5135 5140 5145

Asp Thr Ile Cys Thr His Arg Leu Asp Pro Leu Asn Pro Gly Leu 5150 5160

Asp Arg Glu Gln Leu Tyr Trp Glu Leu Ser Lys Leu Thr Arg Gly 5165 5170 5175

Ile Ile Glu Leu Gly Pro Tyr Leu Leu Asp Arg Gly Ser Leu Tyr 5180 5185 5190

Val Asn Gly Phe Thr His Arg Thr Ser Val Pro Thr Thr Ser Thr 5195 5200 5205

Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe 5210 5220

Ser Leu Pro Ser Pro Ala Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe 5225 5230 5235

Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met 5240 5250

Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 5255 5260 5265

Gln Thr Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu 5270 5280

Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp 5285 5290 5295

Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Leu Asp 5300 5310

Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu Leu 5315 5325

Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr Leu 5330 5340

Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Trp Ile Pro 5345 5350 5355

09-965738substitute.ST25.txt val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly 5370 Ser Gly Thr Pro Ser Leu Pro Ser Ser Pro Thr Thr Ala Gly Pro 5375 5380 5385 Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys 5390 Tyr Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr 5405 5410 5415 5405 Thr Glu Arg Val Leu Gln Ser Leu Leu Gly Pro Met Phe Lys Asn 5425 5430 5420 Thr Ser_Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu_Thr Leu Leu 5440 5445 5435 Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys 5450 5460 Thr His Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln 5465 5470 5475 Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu 5480 5490 Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly 5555 5560 5565 Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly 5585 5590 Page 225

Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr 5605 56Ō0 5610 Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro 5620 5625 Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr 5635 Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser 5645 5650 5655 Leu Tyr Val Asn Gly Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp 5705 5710 5705 Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 5720 5730 Leu Gln Ser Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly 5735 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys 5750 5760 5750 Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His Arg Val 5765 5770 5775 Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp Glu 5780 5785 5790 Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 5800 Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr 5810 5820 Ser Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu 5835 5830 Page 226

Gly	Thr 5840	Ser	Gly	Thr	Pro	Ser 5845	Ser	Leu	Pro	Ser	Pro 5850	Thr	Ser	Αla
Gly	Pro 5855	Leu	Leu	val	Pro	Phe 5860	Thr	Leu	Asn	Phe	Thr 5865	Ile	Thr	Asn
Leu	G1n 5870		Glu	Glu	Asp	Met 5875	His	His	Pro	Gly	Ser 5880	Arg	Lys	Phe
Asn	Thr 5885	Thr	Glu	Arg	Val	Leu 5890	Gln	Gly	Leu	Leu	G]y 5895	Pro	Met	Phe
Lys	Asn 5900	Thr	Ser	val	Gly	Leu 5905	Leu	Tyr	Ser	Glу	Cys 5910	Arg	Leu	Thr
Leu	Leu 5915	Arg	Pro	Glu	Lys	Asn 5920	Gly	Ala	Ala	Thr	Gly 5925	Met	Asp	Ala
Ile	Cys 5930	Thr	His	Arg	Leu	Asp 5935	Pro	Lys	Ser	Pro	Gly 5940	Leu	Asp	Arg
Glu	xaa 5945	Leu	Tyr	Trp	Glu	Leu 5950	Ser	Xaa	Leu	Thr	Xaa 5955	xaa	Ile	xaa
Glu	Leu 5960	Gly	Pro	Tyr	xaa	Leu 5965	Asp	Arg	xaa	Ser	Leu 5970	Туг	٧a٦	Asn
Gly	Phe 5975	Xaa	Xaa	xaa	Xaa	xaa 5980	Xaa	Xaa	Xaa	Thr	Ser 5985	Thr	Pro	Gly
Thr	Ser 5990	Xaa	٧a٦	Xaa	Leu	Xaa 5995	Thr	Ser	Gly	Thr	Pro 6000	Xaa	xaa	Xaa
Pro	xaa 6005	Xaa	Thr	xaa	xaa	xaa 6010	Pro	Leu	Leu	Xaa	Pro 6015	Phe	Thr	Leu
Asn	Phe 6020	Thr	Ile	Thr	Asn	Leu 6025	Xaa	туг	Glu	Glu	xaa 6030	Met	xaa	Xaa
Pro	G]y 6035	Ser	Arg	Lys	Phe	Asn 6040	Thr	Thr	Glu	Arg	va1 6045	Leu	Gln	Gly
Leu	Leu 6050	Lys	Pro	Leu	Phe	Arg 6055	Asn	Ser	ser	Leu	G]u 6060	Tyr	Leu	Tyr
Ser	Gly	Cys	Arg	Leu	Ala	Ser	Leu	Arg	Pro Page	Glu e 227	Lys 7	Asp	Ser	Ser

	Met 6080	Ala	val	Asp	Ala	Ile 6085	Cys	Thr	His	Arg	Pro 6090	Asp	Pro	Glu
Asp	Leu 6095	Gly	Leu	Asp	Arg	Glu 6100	Arg	Leu	Tyr	Тгр	Glu 6105	Leu	Ser	Asn
Leu	Thr 6110	Asn	Gly	Ile	Gln	Glu 6115	Leu	Glу	Pro	Tyr	Thr 6120	Leu	Asp	Arg
Asn	Ser 6125	Leu	Tyr	val	Asn	Gly 6130	Phe	Thr	His	Arg	ser 6135	Ser	Met	Pro
Thr	Thr 6140	Ser	Thr	Pro	Gly	Thr 6145	Ser	Thr	٧al	Asp	Val 6150	Gly	Thr	Ser
Gly	Thr 6155	Pro	Ser	Ser	Ser	Pro 6160	Ser	Pro	Thr	Thr	Ala 6165	Gly	Pro	Leu
Leu	Ile 6170	Pro	Phe	Thr	Leu	Asn 6175	Phe	Thr	Ile	Thr	Asn 6180	Leu	Gln	Tyr
Gly	Glu 6185	Asp	Met	Gly	ніѕ	Pro 6190	Gly	Ser	Arg	Lys	Phe 6195	Asn	Thr	Thr
Glu	Arg 6200	٧a٦	Leu	Gln	Gly	Leu 6205	Leu	Gly	Pro	Ile	Phe 6210	Lys	Asn	Thr
Ser	Val 6215	Gly	Pro	Leu	Tyr	ser 6220	Gly	Cys	Arg	Leu	Thr 6225	Ser	Leu	Arg
Ser	G]u 6230	Lys	Asp	Gly	Аlа	Ala 6235	Thr	Gly	۷al	Asp	Ala 6240	Ile	Cys	Ile
His	ніs 6245	Leu	Asp	Pro	Lys	ser 6250	Pro	Gly	Leu	Asn	Arg 6255	Glu	Arg	Leu
туr	Trp 6260	Glu	Leu	Ser	Gln	Leu 6265	Thr	Asn	Gly	Ile	Lys 6270	Glu	Leu	Gly
Pro	туг 6275	Thr	Leu	Asp	Arg	Asn 6280	Ser	Leu	Tyr	val	Asn 6285	Gly	Phe	Thr
ніѕ	Arg 6290	Thr	Ser	val	Pro	Thr 6295	Thr	Ser	Thr	Pro	Gly 6300	Thr	Ser	Thr

09-965738substitute.ST25.txt Val Asp Leu Gly Thr Ser Gly Thr Pro Phe Ser Leu Pro Ser Pro 6310 Ala Thr Ala Gly Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Thr Leu Leu Gly 6355 Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys 6365 6370 6375 6370 6365 Arg Leu Thr Leu Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly 6385 6380 val Asp_ Ala Ile Cys Thr His Arg Leu Asp Pro Lys_ Ser Pro Gly 6395 6400 Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu 6425 6430 6430 Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro 6460 Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa 6495 Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 6500 6510 Leu Gln Gly Leu Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly 6515 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Lys Lys 6530 6540 6530

Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp Glu 6570 6565 Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Arg Ser 6590 6595 6600 Ser Val Pro Thr Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu 6610 Glu Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro 6625 Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu Arg Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe 6675 6665 6670 Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr 6680 Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr 6695 6700 6705 6695 Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg 6710 6720 Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr 6725 6730 6735 Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val Asp 6740 6745 Gly Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Pro Leu 6775 6780 Page 230

Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu 6785 Asn Phe Thr Ile Thr Asp Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 6825 6820 Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr 6835 6840 6830 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala 6845 6850 6855 Ala Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr 6870 6860 Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg 6900 Asp Ser Leu Tyr Val Asn Gly Phe Asn Pro Trp Ser Ser Val Pro 6910 Thr Thr Ser Thr Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Lys Tyr 6955 Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe Asn Thr Thr 6975 6965 6970 Glu Arg Val Leu Gln Ser Leu His Gly Pro Met Phe Lys Asn Thr 6985 6980 Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg 6995 7000 7005 Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Page 231

His Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Xaa Leu 7025 7030 7035 Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly 7040 7050 Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa 7055 7060 7065 Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr Ser Xaa 7070 7080 Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa 7130 7140 Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys 7145 7150 7155 Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa 7160 7170 Val Asp Xaa Xaa Cys Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly 7175 7180 7185 Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Asn 7190 7195 7200 Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu 7205 7210 7215 Tyr Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser 7220 7225 7230

Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro

09-965738substitute.ST25.txt Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp 7265 7270 7275 Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val 7280 7285 7290 Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 7310 7320 7310 Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Leu 7325 7330 7335 Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu 7340 7350 7350 Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa 7355 7360 7365 Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa 7370 7380 xaa xaa xaa xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe 7440 7435 Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe 7445 7450 Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr 7460 7470 7460 Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa 7480 7475

Xaa Cys Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg 7490 7495 7500 Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa 7505 7510 7515 Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe His Pro Arg Ser Ser Val Pro Thr Thr Ser Thr Pro Gly 7535 7540 7545 Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser Leu Pro Gly His Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 7595 7600 7605 Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr 7610 7615 7620 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asn Gly Ala 7625 7630 7635 Ala Thr Gly Met Asp Ala Ile Cys Ser His Arg Leu Asp Pro Lys 7640 7645 7650 Ser Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg 7670 7680 7670 7680 Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ser Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser 7705 Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Gly Thr 7720 Page 234

Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg 7775 7780 7785 7785 Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa 7790 Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu 7805 7810 7815 Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly 7820 Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Thr 7840 His Gln Asn Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr 7855 Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr 7910 7915 7920 Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys 7925 Arg Leu Thr Leu Leu Arg Pro Glu Lys Gln Glu Ala Ala Thr Gly 7940 7945 Val Asp Thr Ile Cys Thr His Arg Val Asp Pro Ile Gly Pro Gly Page 235

Leu	Asp 7970	Arg	Glu	Xaa	Leu	Tyr 7975	Trp	Glu	Leu	Ser	Xaa 7980	Leu	Thr	Xaa
xaa	11e 7985	xaa	Glu	Leu	Gly	Pro 7990	Tyr	xaa	Leu	Asp	Arg 7995	Xaa	Ser	Leu
Tyr	Val 8000	Asn	G]y	Phe	xaa	xaa 8005	Xaa	xaa	Xaa	xaa	xaa 8010	xaa	Thr	Ser
Thr	Pro 8015	Gly	Thr	Ser	xaa	Val 8020	xaa	Leu	xaa	Thr	Ser 8025	Gly	Thr	Pro
xaa	xaa 8030	Xaa	Pro	xaa	Xaa	Thr 8035	Xaa	xaa	Xaa	Pro	Leu 8040	Leu	xaa	Pro
Phe	Thr 8045	Leu	Asn	Phe	Thr	Ile 8050	Thr	Asn	Leu	Xaa	Tyr 8055	Glu	Glu	Xaa
Met	xaa 8060	Xaa	Pro	Gly	Ser	Arg 8065	Lys	Phe	Asn	Thr	Thr 8070	Glu	Arg	val
Leu	Gln 8075	Gly	Leu	Leu	xaa	Pro 8080	Xaa	Phe	Lys	xaa	Thr 8085	Ser	٧al	Gly
Xaa	Leu 8090		Ser	Gly	Cys	Arg 8095	Leu	Thr	Leu	Leu	Arg 8100	Xaa	Glu	Lys
xaa	Xaa 8105	Ala	Ala	Thr	Xaa	Val 8110	Asp	Xaa	Xaa	Cys	Xaa 8115	Xaa	xaa	Xaa
Asp	Pro 8120	Xaa	Xaa	Pro	Gly	Leu 8125	Asp	Arg	Glu	Xaa	Leu 8130	Tyr	Trp	Glu
Leu	Ser 8135	Xaa	Leu	Thr	Xaa	Xaa 8140		Xaa	Glu	Leu	Gly 8145	Pro	Tyr	xaa
Leu	Asp 8150	Arg	xaa	Ser	Leu	Tyr 8155	Val	Asn	Gly	Phe	Thr 8160	His	Arg	Ser
Ser	Val 8165	Pro	Thr	Thr	Ser	Ser 8170	Pro	Gly	Thr	Ser	Thr 8175	Val	His	Leu
Ala	Thr 8180	Ser	Gly	Thr	Pro	Ser 8185	Ser	Leu	Pro	Gly	His 8190	Thr	Ala	Pro

09-965738substitute.ST25.txt Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn 8205 Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe 8225 8230 8235 Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr 8240 8250 8240 Leu Leu Arg Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala 8255 8260 8265 8255 Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg 8270 8280 8270 Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa 8290 8285 Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly 8315 8320 8325 Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa 8330 Pro Xaa Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly 8375 8380 8385 Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr 8395 8400 8390 Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala 8405 8410 Ala Thr xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa 8420 8425 8430

xaa	Pro 8435	Gly	Leu	Asp	Arg	Glu 8440	Xaa	Leu	Tyr	Trp	Glu 8445	Leu	Ser	xaa
Leu	Thr 8450	xaa	Xaa	Ile	Xaa	Glu 8455	Leu	Gly	Pro	Tyr	xaa 8460	Leu	Asp	Arg
xaa	ser 8465	Leu	туг	val	Asn	Gly 8470	Phe	Thr	His	Arg	Thr 8475	Ser	Val	Pro
Thr	Thr 8480	Ser	Thr	Pro	Gly	Thr 8485	Ser	Thr	val	His	Leu 8490	Ala	Thr	Ser
Gly	Thr 8495	Pro	Ser	Ser	Leu	Pro 8500	Gly	His	Thr	Ala	Pro 8505	val	Pro	Leu
Leu	Ile 8510		Phe	Thr	Leu	Asn 8515	Phe	Thr	Ile	Thr	Asn 8520	Leu	Gln	Tyr
Glu	G1u 8525	Asp	Met	His	Arg	Pro 8530	Gly	Ser	Arg	Lys	Phe 8535	Asn	Thr	Thr
Glu	Arg 8540	val	Leu	Gln	Gly	Leu 8545	Leu	Ser	Pro	Ile	Phe 8550	Lys	Asn	Ser
Ser	Val 8555	Gly	Pro	Leu	Туг	ser 8560	Gly	Cys	Arg	Leu	Thr 8565	Ser	Leu	Arg
Pro	Glu 8570	Lys	Asp	Gly	Ala	Ala 8575	Thr	Gly	Met	Asp	Ala 8580	val	Cys	Leu
Tyr	ніs 8585	Pro	Asn	Pro	Lys	Arg 8590	Pro	Gly	Leu	Asp	Arg 8595	Glu	Gln	Leu
Tyr	Cys 8600	Glu	Leu	Ser	Gln	Leu 8605	Thr	His	Asn	Ile	Thr 8610	Glu	Leu	Gly
Pro	туг 8615	Ser	Leu	Asp	Arg	Asp 8620	Ser	Leu	Tyr	val	Asn 8625	Gly	Phe	Thr
His	G]n 8630	Asn	Ser	val	Pro	Thr 8635	Thr	Ser	Thr	Pro	Gly 8640	Thr	Ser	Thr
٧a٦	Tyr 8645	Тгр	Ala	Thr	Thr	Gly 8650	Thr	Pro	Ser	Ser	Phe 8655	Pro	Gly	His
Thr	Xaa 8660		xaa	Pro	Leu	Leu 8665	xaa	Pro		Thr e 23	Leu 8670 8	Asn	Phe	Thr

Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met Xaa Xaa Pro Gly Ser 8680 Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa 8720 Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly 8735 Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa 8750 8760 Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu 8765 8770 8775 8765 Tyr Val Asn Gly Phe Thr His Trp Ser Ser Gly Leu Thr Thr Ser Thr Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro 8795 8800 8805 Ser Pro Val Pro Ser Pro Thr Thr Ala Gly Pro Leu Leu Val Pro 8815 8820 Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys Asn Thr Ser Val Gly 8865 8855 8860 Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys 8870 8880 Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His Arg Val 8890 8895 8885 Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Page 239

Leu Ser 891	xaa .5	Leu	Thr	Xaa	Xaa 8920	Ile	xaa	Glu	Leu	G]y 8925	Pro	Tyr	Xaa
Leu Asp 893		xaa	Ser	Leu	Tyr 8935	val	Asn	G1y	Phe	xaa 8940	xaa	xaa	Xaa
Xaa Xaa 894		Xaa	Thr	Ser	Thr 8950		Gly	Thr	Ser	Xaa 8955	val	Xaa	Leu
Xaa Thi 896		Gly	Thr	Pro	Xaa 8965	xaa	Xaa	Pro	xaa	Xaa 8970	Thr	xaa	xaa
Xaa Pro 897		Leu	Xaa	Pro	Phe 8980	Thr	Leu	Asn	Phe	Thr 8985	Ile	Thr	Asn
Leu Xaa 899		Glu	Glu	Xaa	Met 8995	Xaa	xaa	Pro	Gly	ser 9000	Arg	Lys	Phe
Asn Thi 900		Glu	Arg	val	Leu 9010	Gln	Gly`	Leu	Leu	Xaa 9015	Pro	xaa	Phe
Lys Xaa 902		Ser	٧a٦	Gly	xaa 9025	Leu	Tyr	Ser	Gly	Cys 9030	Arg	Leu	Thr
Leu Lei 903		Xaa	Glu	Lys	Xaa 9040	Xaa	Ala	Ala	Thr	Xaa 9045	val	Asp	Xaa
Xaa Cys 905		Xaa	Xaa	Xaa	Asp 9055	Pro	xaa	xaa	Pro	G]y 9060	Leu	Asp	Arg
Glu Xaa 906		Tyr	Тгр	Glu	Leu 9070	ser	xaa	Leu	Thr	xaa 9075	Xaa	Ile	xaa
Glu Lei 908		Pro	Туг	Xaa	Leu 9085	Asp	Arg	xaa	ser	Leu 9090	Туг	٧a٦	Asn
Gly Phe 909		His	Arg	ser	Phe 9100	Gly	Leu	Thr	Thr	ser 9105	Thr	Pro	Trp
Thr Sei 913		val	Asp	Leu	Gly 9115	Thr	Ser	Gly	Thr	Pro 9120	Ser	Pro	val
Pro Sei 912		Thr	Thr	Ala	Gly 9130	Pro	Leu	Leu	٧a٦	Pro 9135	Phe	Thr	Leu

09-965738substitute.ST25.txt Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His Arg 9145 9150 Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Thr Pro Leu Phe Arg Asn Thr Ser Val Ser Ser Leu Tyr 9170 Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Ala 9185 9190 9195 9190 Ala Thr Arg Val Asp Ala Val Cys Thr His Arg Pro Asp Pro Lys 9200 9210 9200 Ser Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa 9225 9215 Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg 9230 Xaa Ser Leu Tyr Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa 9245 9250 9255 Xaa Thr Ser Thr Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser 9260 9265 9270 Gly Thr Pro Xaa Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu 9285 9275 Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr 9300 Glu Glu Xaa Met Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser val Gly Xaa Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg 9340 9345 9335 Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa 9350 Xaa Xaa Xaa Asp Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu 9365 9370 9375

Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly 9390 9380 9385 Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Thr 9395 9400 9405 His Trp Ile Pro Val Pro Thr Ser Ser Thr Pro Gly Thr Ser Thr 9415 Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr 9425 9430 9435 Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro 9470 9480 Ile Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val 9510 9505 9500 Asp Ala Ile Cys Ile His His Leu Asp Pro Lys Ser Pro Gly Leu 9525 9515 Asp Arg Glu Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa 9530 9540 Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr 9545 9550 9555 9545 Val Asn Gly Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr 9570 9560 Pro Gly Thr Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa 9575 9580 9585 Xaa Xaa Pro Xaa Xaa Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met 9610 Page 242

Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Xaa Pro Xaa Phe Lys Xaa Thr Ser Val Gly Xaa 9645 Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Xaa Glu Lys Xaa Xaa Ala Ala Thr Xaa Val Asp Xaa Xaa Cys Xaa Xaa Xaa Xaa Asp 9665 9670 9675 Pro Xaa Xaa Pro Gly Leu Asp Arg Glu Xaa Leu Tyr Trp Glu Leu 9680 9685 9690 9680 Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Leu Gly Pro Tyr Xaa Leu 9705 9695 Asp Arg Xaa Ser Leu Tyr Val Asn Gly Phe Thr His Gln Thr Phe Ala Pro Asn Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly 9725 9730 9735 Thr Ser Gly Thr Pro Ser Ser Leu Pro Ser Pro Thr Ser Ala Gly 9740 Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe Asn 9775 Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly Cys Arg Leu Thr Leu 9810 9805 9800 Leu Arg Pro Glu Lys Asn Gly Ala Ala Thr Arg Val Asp Ala Val 9815 Cys Thr His Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu 9835 9840 9830 Xaa Leu Tyr Trp Glu Leu Ser Xaa Leu Thr Xaa Xaa Ile Xaa Glu Page 243

Leu Gly Pro Tyr Xaa Leu Asp Arg Xaa Ser Leu Tyr Val Asn Gly 9860 9870

Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa Thr Ser Thr Pro Gly Thr 9875 9880 9885

Ser Xaa Val Xaa Leu Xaa Thr Ser Gly Thr Pro Xaa Xaa Xaa Pro 9890 9895 9900

Xaa Xaa Thr Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn 9905 9910 9915

Phe Thr Ile Thr Asn Leu His Tyr Glu Glu Asn Met Gln His Pro 9920 9925 9930

Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 9935 9940 9945

Leu Arg Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser 9950 9955 9960

Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys His Gly Ala Ala 9965 9970 9975

Thr Gly Val Asp Ala Ile Cys Thr Leu Arg Leu Asp Pro Thr Gly 9980 9985 9990

Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Gln Leu 9995 10000 10005

Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp 10010 10015 10020

Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser Val Pro Thr 10025 10030 10035

Thr Ser Ile Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly 10040 10045 10050

Thr Pro Ala Ser Leu Pro Gly His Thr Ala Pro Gly Pro Leu Leu 10055 10060 10065

Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu 10070 10075 10080

09-965738substitute.ST25.txt Val Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu 10085 10090 10095 Arg Val Leu Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser 10105 Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro 10115 10120 10125 Glu Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His 10130 10135 10140 Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr 10145 10150 10155 Trp Glu Leu Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro 10170 10160 Tyr Leu Leu Asp Arg Gly Ser Leu Tyr Val Asn Gly Phe Thr His 10175 10180 10185Arg Asn Phe Val Pro Ile Thr Ser Thr Pro Gly Thr Ser Thr Val 10190 10200 His Leu Gly Thr Ser Glu Thr Pro Ser Ser Leu Pro Arg Pro Ile 10205 10210 10215Val Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile 10220 10225 10230 Thr Asn Leu Gln Tyr Glu Glu Ala Met Arg His Pro Gly Ser Arg 10240 Lys Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 10250 10255 1026 Leu Arg Pro Leu Phe Lys Asn Thr Ser Ile Gly Pro Leu Tyr Ser Ser Cys Arg 10265 10270 10275 10270 Leu Thr Leu Leu Arg Pro Glu Lys Asp Lys Ala Ala Thr Arg Val 10280 10285 10290 Asp Ala Ile Cys Thr His His Pro Asp Pro Gln Ser Pro Gly Leu 10295 Asn Arg Glu Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly 10310 10320 10310

Ile Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr 10330 10325 Val Asp Gly Phe Thr His Trp Ser Pro Ile Pro Thr Thr Ser Thr 10340 10345 10350Pro Gly Thr Ser Ile Val Asn Leu Gly Thr Ser Gly Ile Pro Pro 10355 10360 10365 Ser Leu Pro Glu Thr Thr Xaa Xaa Xaa Pro Leu Leu Xaa Pro Phe 10370 10375 10380 Thr Leu Asn Phe Thr Ile Thr Asn Leu Xaa Tyr Glu Glu Xaa Met 10385 10390 10395 Xaa Xaa Pro Gly Ser Arg Lys Phe Asn Thr Thr Glu Arg Val Leu 10405 10410 Gln Gly Leu Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro 10420 Leu Tyr Ser Gly Cys Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp 10430 10435 10440 Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His Arg Pro Asp 10445 10450 10455 Pro Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp Glu Leu 10460 10465 10470 10460 Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr Leu 10475 10480 10485 10475 Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr Gln Arg Ser Ser 10490 10495 10500 Val Pro Thr Thr Ser Thr Pro Gly Thr Phe Thr Val Gln Pro Glu 10505 10510 10515 Thr Ser Glu Thr Pro Ser Ser Leu Pro Gly Pro Thr Ala Thr Gly 10525 Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu 10535 10545 10540 Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe Asn 10550 10560 Page 246

Thr	Thr 10565	Glu	Arg	۷al	Leu	Gln 10570	Gly	Leu	Leu	Met	Pro 10575	Leu	Phe	Lys
Asn	Thr 10580	Ser	Val	Ser	Ser	Leu 10585	Tyr	Ser	Gly	Cys	Arg 10590	Leu	Thr	Leu
Leu	Arg 10595	Pro	Glu	Lys	Asp	Gly 10600	Ala	Ala	Thr	Arg	val 10605	Asp	Ala	val
Cys	Thr 10610	His	Arg	Pro	Asp	Pro 10615	Lys	Ser	Pro	Glу	Leu 10620	Asp	Arg	Glu
Arg	Leu 10625	Tyr	Trp	Lys	Leu	Ser 10630	Gln	Leu	Thr	His	Gly 10635	Ile	Thr	Glu
Leu	Gly 10640	Pro	Tyr	Thr	Leu	Asp 10645	Arg	His	Ser	Leu	Туг 10650	val	Asn	Gly
Phe	Thr 10655	нis	G∏n	Ser	Ser	Met 10660	Thr	Thr	Thr	Arg	Thr 10665	Pro	Asp	Thr
Ser	Thr 10670	Met	His	Leu	Ala	Thr 10675	Ser	Arg	Thr	Pro	Ala 10680	Ser	Leu	Ser
Gly	Pro 10685		Thr	Ala	Ser	Pro 10690	Leu	Leu	val	Leu	Phe 10695	Thr	Ile	Asn
Phe	Thr 10700	Ile	Thr	Asn	Leu	Arg 10705	Tyr	Glu	Glu	Asn	Met 10710	His	His	Pro
Gly	Ser 10715	Arg	Lys	Phe	Asn	Thr 10720	Thr	Glu	Arg	∨al	Leu 10725	Gln	Gly	Leu
Leu	Arg 10730	Pro	val	Phe	Lys	Asn 10735	Thr	Ser	val	Gly	Pro 10740	Leu	Tyr	Ser
Glу	Cys 10745	Arg	Leu	Thr	Leu	Leu 10750	Arg	Pro	Lys	Lys	Asp 10755	Gly	Ala	Ala
Thr	Lys 10760	val	Asp	Αla	Ile	Cys 10765	Thr	туг	Arg	Pro	Asp 10770	Pro	Lys	Ser
Pro	Gly 10775	Leu	Asp	Arg	Glu	Gln 10780	Leu	Туг	Тгр	Glu	Leu 10785	Ser	Gln	Leu
Thr	His	Ser	Ile	Thr	Glu	Leu	Gly	Pro Pa	Tyr age 2	Thr 247	Gln	Asp	Arg	Asp

Ser	Leu 10805	Tyr	Asn	val	Glу	Phe 10810	Thr	Gln	Arg	Ser	Ser 10815	val	Pro	Thr
Thr	ser 10820		Pro	Gly	Thr	Pro 10825	Thr	val	Asp	Leu	Gly 10830	Thr	Ser	Gly
Thr	Pro 10835		Ser	Lys	Pro	Gly 10840	Pro	Ser	Αla	Ala	Ser 10845	Pro	Leu	Leu
Val	Leu 10850	Phe	Thr	Leu	Asn	Gly 10855	Thr	Ile	Thr	Asn	Leu 10860	Arg	Tyr	Glu
Glu	Asn 10865	Met	Gln	His	Pro	Gly 10870	Ser	Arg	Lys	Phe	Asn 10875	Thr	Thr	Glu
Arg	Val 10880	Leu	Gln	Gly	Leu	Leu 10885	Arg	Ser	Leu	Phe	Lys 10890	Ser	Thr	Ser
val	Gly 10895	Pro	Leu	Tyr	Ser	Gly 10900	Cys	Arg	Leu	Thr	Leu 10905	Leu	Arg	Pro
Glu	Lys 10910	Asp	Gly	Thr	Ala	Thr 10915	Gly	val	Asp	Ala	Ile 10920	Cys	Thr	His
His	Pro 10925	Asp	Pro	Lys	ser	Pro 10930	Arg	Leu	Asp	Arg	Glu 10935	Gln	Leu	Tyr
Trp	Glu 10940		Ser	Gln	Leu	Thr 10945	His	Asn	Ile	Thr	Glu 10950	Leu	Gly	His
Tyr	Ala 10955	Leu	Asp	Asn	Asp	Ser 10960	Leu	Phe	val	Asn	Gly 10965	Phe	Thr	His
Arg	Ser 10970	Ser	۷al	Ser	Thr	Thr 10975	Ser	Thr	Pro	Gly	Thr 10980	Pro	Thr	val
Tyr	Leu 10985		Αla	Ser	Lys	Thr 10990	Pro	Ala	Ser	Ile	Phe 10995	Gly	Pro	Ser
Ala	Ala 11000		ніѕ	Leu	Leu	Ile 11005	Leu	Phe	Thr	Leu	Asn 11010	Phe	Thr	Ile
Thr	Asn 11015		Arg	Туг	Glu	Glu 11020	Asn	Met	Trp	Pro	Gly 11025	Ser	Arg	Lys

09-965738substitute.ST25.txt Phe Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Arg Pro Leu 11035 11040 Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Ser Arg Leu Thr Leu Leu Arg Pro Glu Lys Asp Gly Glu Ala Thr Gly Val Asp 11060 11065 11070 Ala Ile Cys Thr His Arg Pro Asp Pro Thr Gly Pro Gly Leu Asp 11075 11080 11085 Arg Glu Gln Leu Tyr Leu Glu Leu Ser Gln Leu Thr His Ser Ile 11090 11095 11100 Thr Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asp Ser Leu Tyr Val 11110 11115 11105 Asn Gly Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Thr Gly 11125 11130 Val Val Ser Glu Glu Pro Phe Thr Leu Asn Phe Thr Ile Asn Asn 11135 11140 11145 Leu Arg $\,$ Tyr Met Ala Asp Met $\,$ Gly Gln Pro Gly Ser $\,$ Leu Lys Phe $\,$ 11150 $\,$ 11160 Asn Ile Thr Asp Asn Val Met Lys His Leu Leu Ser Pro Leu Phe 11170 Gln Arg Ser Ser Leu Gly Ala Arg Tyr Thr Gly Cys Arg Val Ile 11180 11185 11190 Ala Leu Arg Ser Val Lys Asn Gly Ala Glu Thr Arg Val Asp Leu 11200 Leu Cys Thr Tyr Leu Gln Pro Leu Ser Gly Pro Gly Leu Pro Ile 11210 11220 11215 Lys Gln Val Phe His Glu Leu Ser Gln Gln Thr His Gly Ile Thr 11225 11230 11235 Arg Leu Gly Pro Tyr Ser Leu Asp Lys Asp Ser Leu Tyr Leu Asn 11245 11240 Gly Tyr Asn Glu Pro Gly Leu Asp Glu Pro Pro Thr Thr Pro Lys 11255 11260 11265

Pro Ala Thr Thr Phe Leu Pro Pro Leu Ser Glu Ala Thr Thr Ala Met Gly Tyr His Leu Lys Thr Leu Thr Leu Asn Phe Thr Ile Ser 11285 11290 11295 Asn Leu Gln Tyr Ser Pro Asp Met Gly Lys Gly Ser Ala Thr Phe 11300 11305 11310Asn Ser Thr Glu Gly Val Leu Gln His Leu Leu Arg Pro Leu Phe 11315 11320 11325 Gln Lys Ser Ser Met Gly Pro Phe Tyr Leu Gly Cys Gln Leu Ile 11330 11335 11340 Ser Leu Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr 11345 11350 11355 Thr Cys Thr Tyr His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile 11360 11365 11370 Gln Gln Leu Tyr Trp Glu Leu Ser Gln Leu Thr His Gly Val Thr 11375 11380 11385 Gln Leu Gly Phe Tyr Val Leu Asp Arg Asp Ser Leu Phe Ile Asn 11390 . 11395 11400 Gly Tyr Ala Pro Gln Asn Leu Ser Ile Arg Gly Glu Tyr Gln Ile 11405 11410 11415 Asn Phe His Ile Val Asn Trp Asn Leu Ser Asn Pro Asp Pro Thr 11420 11425 11430 Ser Ser Glu Tyr Ile Thr Leu Leu Arg Asp Ile Gln Asp Lys Val 11435 11440 11445 Thr Thr Leu Tyr Lys Gly Ser Gln Leu His Asp Thr Phe Arg Phe 11450 11460 Cys Leu Val Thr Asn Leu Thr Met Asp Ser Val Leu Val Thr Val 11470 Lys Ala Leu Phe Ser Ser Asn Leu Asp Pro Ser Leu Val Glu Gln 11480 11490 Val Phe Leu Asp Lys Thr Leu Asn Ala Ser Phe His Trp Leu Gly 11500 11505 Page 250

Ser Thr Tyr Gln Leu Val Asp Ile His Val Thr Glu Met Glu Ser Ser Val Tyr Gln Pro Thr Ser Ser Ser Ser Thr Gln His Phe Tyr 11525 11530 11535 11530 Leu Asn Phe Thr Ile Thr Asn Leu Pro Tyr Ser Gln Asp Lys Ala 11540 11550 11545 Gln Pro Gly Thr Thr Asn Tyr Gln Arg Asn Lys Arg Asn Ile Glu 11555 11560 11565 Asp Ala Leu Asn Gln Leu Phe Arg Asn Ser Ser Ile Lys Ser Tyr 11570 11580 11570 Phe Ser Asp Cys Gln Val Ser Thr Phe Arg Ser Val Pro Asn Arg 11590 His His Thr Gly Val Asp Ser Leu Cys Asn Phe Ser Pro Leu Ala 11600 11605 11610 Arg Arg Val Asp Arg Val Ala Ile Tyr Glu Glu Phe Leu Arg Met 11615 11620 11625 Thr Arg Asn Gly Thr Gln Leu Gln Asn Phe Thr Leu Asp Arg Ser 11630 11640 Ser Val Leu Val Asp Gly Tyr Ser Pro Asn Arg Asn Glu Pro Leu 11650 Thr Gly Asn Ser Asp Leu Pro Phe Trp Ala Val Ile Leu Ile Gly 11665 Leu Ala Gly Leu Leu Gly Leu Ile Thr Cys Leu Ile Cys Gly Val 11675 11680 11685Leu Val Thr Thr Arg Arg Arg Lys Lys Glu Gly Glu Tyr Asn Val 11690 11695 11700 Gln Gln Cys Pro Gly Tyr Tyr Gln Ser His Leu Asp Leu Glu 11705 11710 11715 Asp Leu Gln 11720

<210> 163

<211> 156

<212> PRT

<213> Homo sapiens

<400> 163

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30

Asn Ala Thr Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys 35 40 45

Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu 50 60

Arg Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu 65 70 75 80

Tyr His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr 85 90 95

Trp Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr 100 105 110

Ser Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly Phe Thr His Gln Asn 115 120 125

Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Tyr Trp Ala 130 135 140

Thr Thr Gly Thr Pro Ser Ser Phe Pro Gly His Thr 145 150 155

<210> 164

<211> 42

<212> PRT

<213> Homo sapiens

<400> 164

Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Ala Thr Glu Arg Glu Leu Gln Gly Leu 35 40

```
<210> 165
<211>
      42
<212>
      PRT
<213>
       Homo sapiens
<400>
       165
Thr Ala Val Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr
Asn Leu Gln Tyr Gly Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
<210>
     166
<211>
       42
<212>
       PRT
<213>
       Homo sapiens
<400>
       166
Val Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15
Asn Leu Gln Tyr Glu Glu Ala Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
<210> 167
<211>
       42
<212>
       PRT
<213>
       Homo sapiens
<400> 167
Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15
Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30
Ser Thr Thr Glu Arg Val Leu Gln Gly Leu
<210>
       168
<211>
       42
<212>
       PRT
<213>
       Homo sapiens
```

<400> 168

Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

<210> 169

<211> <212> 42

PRT <213> Homo sapiens

<400>

Ala Pro Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 10 15

Asn Leu Gln Tyr Glu Val Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

<210> 170

<211> <212> 42

PRT <213> Homo sapiens

<400> 170

Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

<210> 171

<211> 42

<212> **PRT**

<213> Homo sapiens

<400> 171

Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe 20 25 30

Asn Thr Met Glu Ser Val Leu Gln Gly Leu 35 40

<210> 172

<211> 42 <212> PRT

<213> Homo sapiens

<400> 172

Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Cys Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Glu Glu Asp Met Arg Arg Thr Gly Ser Arg Lys Phe 20 25 30

Asn Thr Met Glu Ser Val Leu Gln Gly Leu 35 40

<210> 173

<211> 42

<212> PRT

<213> Homo sapiens

<400> 173

Ala Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 174

<211> 42

<212> PRT

<213> Homo sapiens

<400> 174

Thr Ala Gly Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

Page 255

```
09-965738substitute.ST25.txt
```

35

<210> 175

<211> 42

<212> PRT <213> Homo sapiens

<400> 175

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Gly Glu Asp Met Gly His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 176

<211> 42

<212> PRT

<213> Homo sapiens

<400> 176

Thr Ala Gly Pro Leu Leu Val Leu Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Lys Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Thr Leu 35 40

<210> 177

<211> 42

<212> PRT

<213> Homo sapiens

<400> 177

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30

Asn Ala Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 178

<211> 42

<212> PRT

```
09-965738substitute.ST25.txt
```

<213> Homo sapiens

<400> 178

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Arg Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 179

<211> 42

<212> PRT

<213> Homo sapiens

<400> 179

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
35 40

<210> 180

<211> 42

<212> PRT <213> Homo sapiens

<400> 180

Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 181

<211> 42

<212> PRT <213> Homo sapiens

<400> 181

Ala Thr Gly Pro Val Leu Leu Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15 Page 257

Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 182

<211> 42

<212> PRT

<213> Homo sapiens

<400> 182

Ala Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 5 10 15

Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

<210> 183

<211> 42

<212> PRT

<213> Homo sapiens

<400> 183

Ser Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Glu Glu Asp Met His His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 184

42

<211> <212> PRT

<213> Homo sapiens

<400> 184

Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr 1 10 15

Asn Gln Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe $20 \hspace{1cm} 25 \hspace{1cm} 30$

```
09-965738substitute.ST25.txt
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
<210>
        185
<211>
<212>
<213>
        42
        PRT
        Homo sapiens
<400>
        185
Thr Ala Ser Pro Leu Leu Val Leu Phe Thr Ile Asn Phe Thr Ile Thr 1 10 15
Asn Leu Arg Tyr Glu Glu Asn Met His His Pro Gly Ser Arg Lys Phe
20 25 30
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
<210>
        186
<211>
<212>
        42
        PRT
<213>
        Homo sapiens
<400>
        186
Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr 1 10 15
Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe 20 25 30
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
<210>
        187
        42
<211>
<212>
        PRT
<213>
        Homo sapiens
<400>
        187
Glu Pro Gly Pro Leu Leu Ile Pro Phe Thr Phe Asn Phe Thr Ile Thr 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Asn Leu Arg Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe 20 \hspace{1cm} 25 \hspace{1cm} 30
Asn Thr Thr Glu Arg Val Leu Gln Gly Leu
         35
<210>
        188
<211> 42
```

```
09-965738substitute.ST25.txt
```

<212> PRT

<213> Homo sapiens

<400> 188

Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 10 15

Asn Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

<210> 189

<211> 42

<212> PRT

<213> Homo sapiens

<400> 189

Ala Pro Val Pro Leu Leu Ile Pro Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asp Leu His Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu

<210> 190

42

<211> <212> PRT

<213> Homo sapiens

<400> 190

Ala Ala Ser Pro Leu Leu Val Leu Phe Thr Leu Asn Gly Thr Ile Thr 1 $5101510151010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101010101$

Asn Leu Arg Tyr Glu Glu Asn Met Gln His Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 191

<211> 42

<212> PRT

<213> Homo sapiens

<400> 191

Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Page 260

Asn Leu Lys Tyr Glu Glu Asp Met His Cys Pro Gly Ser Arg Lys Phe 20 25 30

Asn Thr Thr Glu Arg Val Leu Gln Ser Leu 35 40

5

192 <210>

<211> 41

<212> PRT <213> Homo sapiens

<400> 192

Ala Ala Ser His Leu Leu Ile Leu Phe Thr Leu Asn Phe Thr Ile Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Arg Tyr Glu Glu Asn Met Trp Pro Gly Ser Arg Lys Phe Asn 20 25 30

Thr Thr Glu Arg Val Leu Gln Gly Leu 35 40

<210> 193

<211> 42

<212> PRT

<213> Homo sapiens

<400> 193

Asn Leu Arg Tyr Met Ala Asp Met Gly Gln Pro Gly Ser Leu Lys Phe $20 \hspace{1cm} 25 \hspace{1cm} 30$

Asn Ile Thr Asp Asn Val Met Lys His Leu 35 40

<210> 194

42

<211> <212> PRT

Homo sapiens <213>

<400>

Ala Met Gly Tyr His Leu Lys Thr Leu Thr Leu Asn Phe Thr Ile Ser $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Asn Leu Gln Tyr Ser Pro Asp Met Gly Lys Gly Ser Ala Thr Phe Asn 20 25 30

```
Ser Thr Glu Gly Val Leu Gln His Leu Leu
<210>
        195
        23
<211>
<212>
       PRT
<213> Homo sapiens
<400> 195
Leu Lys Pro Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Ala Ser Leu Arg
20
<210> 196
<211> 23
<212> PRT
<213> Homo sapiens
<400> 196
Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly 1 5 10
Cys Arg Leu Thr Leu Leu Arg
             20
<210> 197
<211> 23
<212> PRT
<213> Homo sapiens
<400> 197
Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 5 10 15
Cys Arg Leu Thr Leu Leu Arg
<210> 198
<211> 23
       198
<212> PRT
<213> Homo sapiens
<400> 198
Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 5 10 15
Cys Arg Leu Thr Leu Leu Arg
```

```
<210> 199
       23
<211>
<212>
       PRT
<213> Homo sapiens
<400> 199
Leu Lys Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Ser 1 5 10 15
Cys Arg Leu Thr Leu Leu Arg
<210> 200
<211> 23
<212> PRT
<213> Homo sapiens
<400> 200
Leu Lys Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Ser Leu Arg
              20
<210>
        201
<211>
        23
<212>
        PRT
<213> Homo sapiens
<400>
      201
Leu Gly Pro Ile Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 10 \ 15
Cys Arg Leu Thr Ser Leu Arg
              20
<210> 202
<211>
        23
<212> PRT
<213>
      Homo sapiens
<400>
        202
Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Leu Leu Arg
              20
<210> 203
```

```
09-965738substitute.ST25.txt
<211> 23
<212> PRT
<213> Homo sapiens
<400> 203
Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Leu Leu Arg
<210>
<211>
        204
        23
<212> PRT
<213> Homo sapiens
<400> 204
Leu Gly Pro Met Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 10 15
Cys Arg Leu Thr Ser Leu Arg
<210> 205
<211> 23
<212> PRT
<213> Homo sapiens
<400> 205
Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly 10 15
Cys Arg Leu Ile Ser Leu Arg
20
<210> 206
<211>
       23
<212>
       PRT
<213>
        Homo sapiens
<400> 206
Leu Gly Pro Leu Phe Lys Asn Ser Ser Val Asp Pro Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Ser Leu Arg
              20
<210>
        207
<211>
        23
<212> PRT
```

<213> Homo sapiens

```
<400> 207
Leu Ser Pro Ile Phe Lys Asn Ser Ser Val Gly Pro Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Ser Leu Arg
              20
<210> 208
<211> 23
<212> PRT
<213> Homo sapiens
<400>
        208
Leu Ser Pro Ile Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 10 15
Cys Arg Leu Thr Leu Leu Arg
              20
<210>
        209
<211> 23
<212> PRT
<213> Homo sapiens
<400> 209
Leu Ser Pro Leu Phe Gln Arg Ser Ser Leu Gly Ala Arg Tyr Thr Gly 1 5 10 15
Cys Arg Val Ile Ala Leu Arg
              20
        210
<210>
        23
<211>
<212>
        PRT
<213>
        Homo sapiens
<400>
        210
Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly 1 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Leu Leu Arg
              20
<210>
        211
<211>
        23
<212>
        PRT
<213>
        Homo sapiens
<400> 211
```

```
09-965738substitute.ST25.txt
Leu Arg Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 5 10 15
Ser Arg Leu Thr Leu Leu Arg
<210> 212
        23
<211>
<212>
        PRT
<213>
        Homo sapiens
<400> 212
Leu Arg Pro Leu Phe Lys Asn Thr Ser Ile Gly Pro Leu Tyr Ser Ser 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Leu Leu Arg
              20
<210>
        213
<211>
        23
<212>
        PRT
<213> Homo sapiens
<400> 213
Leu Arg Pro Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Leu Leu Arg
              20
<210>
        214
<211>
        23
<212> PRT
<213> Homo sapiens
<400> 214
Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Leu Leu Tyr Ser Gly
Cys Arg Leu Thr Leu Leu Arg
              20
<210> 215
<211> 23
<212> PRT
<213> Homo sapiens
<400> 215
Leu Arg Pro Val Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 5 10 15
```

```
Cys Arg Leu Thr Leu Leu Arg
              20
<210>
        216
        23
<211>
<212>
        PRT
<213> Homo sapiens
<400> 216
Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Leu Leu Arg
              20
<210> 217
<211> 23
<212> PRT
<213> Homo sapiens
<400> 217
Leu Arg Ser Leu Phe Lys Ser Thr Ser Val Gly Pro Leu Tyr Ser Gly 1 5 10 15
Cys Arg Leu Thr Ser Leu Arg
<210> 218
<211> 23
<212> PRT
<213> Homo sapiens
<400> 218
Leu Thr Pro Leu Phe Lys Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly 10 15
Cys Arg Leu Thr Leu Leu Arg
<210> 219
<211> 23
<212> PRT
<213> Homo sapiens
<400> 219
Leu Thr Pro Leu Phe Arg Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Cys Arg Leu Thr Leu Leu Arg
              20
```

```
<210> 220
<211> 23
<212> PRT
<213>
       Homo sapiens
<400>
       220
Leu Met Pro Leu Phe Lys Asn Thr Ser Val Ser Ser Leu Tyr Ser Gly
1 10 15
Cys Arg Leu Thr Leu Leu Arg
<210> 221
<211> 22
<212> PRT
<213> Homo sapiens
<400>
       221
Arg Pro Leu Phe Gln Lys Ser Ser Met Gly Pro Phe Tyr Leu Gly Cys 1 \hspace{1cm} 10 \hspace{1cm} 15
Gln Leu Ile Ser Leu Arg
20
<210> 222
<211>
<212>
        58
        PRT
<213>
       Homo sapiens
<400>
Pro Glu Lys Asp Ser Ser Ala Met Ala Val Asp Ala Ile Cys Thr His 1 10 15
Arg Pro Asp Pro Glu Asp Leu Gly Leu Asp Arg Glu Arg Leu Tyr Trp 20 \hspace{1cm} 25 \hspace{1cm} 30
Glu Leu Ser Asn Leu Thr Asn Gly Ile Gln Glu Leu Gly Pro Tyr Thr 35 40 45
Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50
<210> 223
<211> 58
<212> PRT
<213> Homo sapiens
<400> 223
Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His
                                            Page 268
```

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50

5

<210> 224

<211> <212> 58

PRT

<213> Homo sapiens

224 <400>

Pro Lys Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His 1 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Lys Leu Thr Asn Asp Ile Glu Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50

<210> 225

<211> 58

<212> PRT

<213> Homo sapiens

<400> 225

Pro Glu Lys Asp Gly Thr Ala Thr Gly Val Asp Ala Ile Cys Thr His 1 10 15

His Pro Asp Pro Lys Ser Pro Arg Leu Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly His Tyr Ala 35 40 45

Leu Asp Asn Asp Ser Leu Phe Val Asn Gly

<210> 226 <211> 58

<212> PRT

<213> Homo sapiens

<400> 226

Pro Glu Lys Asp Gly Glu Ala Thr Gly Val Asp Ala Ile Cys Thr His 1 10 15

Arg Pro Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Gln Leu Tyr Leu 20 25 30

Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 227

<211> 58

<212> PRT

<213> Homo sapiens

<400> 227

Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr 1 10 15

His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 228

<211> 58

<212> PRT

<213> Homo sapiens

<400> 228

Pro Glu Lys Asp Gly Ala Ala Thr Gly Met Asp Ala Val Cys Leu Tyr 1 10 15

His Pro Asn Pro Lys Arg Pro Gly Leu Asp Arg Glu Gln Leu Tyr Cys 20 25 30

Glu Leu Ser Gln Leu Thr His Asn Ile Thr Glu Leu Gly Pro Tyr Ser 40 45

```
09-965738substitute.ST25.txt
Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50
```

<210> 229 <211> 58

<212> PRT

<213> Homo sapiens

229 <400>

Pro Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Ala Cys Thr Tyr
1 10 15

Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Val Ser Leu Tyr Val Asn Gly 50 55

230 <210>

<211> <212> 58

PRT

<213> Homo sapiens

<400> 230

Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr
1 5 10 15

Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Gln Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 231

<211> 58

<212> PRT

Homo sapiens <213>

<400> 231

Pro Lys Lys Asp Gly Ala Ala Thr Lys Val Asp Ala Ile Cys Thr Tyr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp 20 25 30 Page 271

Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Gln Asp Arg Asp Ser Leu Tyr Asn Val Gly 50

<210> 232 <211> 58

<212> PRT Homo sapiens <213>

<400> 232

Pro Glu Lys Asp Gly Ala Ala Thr Arg Val Asp Ala Val Cys Thr His 1 10 15

Arg Pro Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp 20 25 30

Lys Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg His Ser Leu Tyr Val Asn Gly 50 55

<210> 233 <211> 58 <212> PRT <213> Homo sapiens

<400> 233

Pro Glu Lys Asp Gly Val Ala Thr Arg Val Asp Ala Ile Cys Thr His 1 10 15

Arg Pro Asp Pro Lys Ile Pro Gly Leu Asp Arg Gln Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr His Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 234

<211> 58

<212> PRT <213> Homo sapiens

<400> 234

09-965738substitute.ST25.txt Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Ile His 1 10 15

His Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Arg Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50

<210> 235 <211> 58 <212> PRT

<213> Homo sapiens

<400> 235

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His 1 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50

<210> 236 <211> 58 <212> PRT

<213> Homo sapiens

<400> 236

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His 1 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50 55

<210> 237

<211> 58

<212> **PRT**

Homo sapiens

<400> 237

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His 1 10 15

Arg Val Asp Pro Lys Ser Pro Gly Val Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50

<210> 238

<211> 58

PRT

<212> <213> Homo sapiens

<400> 238

Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr His 1 10 15

His Leu Asn Pro Gln Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Gln Leu Ser Gln Met Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50

239 <210>

<211> 58

<212> PRT

<213> Homo sapiens

<400>

Pro Glu Lys Arg Gly Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His 1 10 15

Arg Leu Asp Pro Leu Asn Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Lys Leu Thr Arg Gly Ile Ile Glu Leu Gly Pro Tyr Leu 35 40 45

Leu Asp Arg Gly Ser Leu Tyr Val Asn Gly 50

<210> 240

<211> 58

<212> PRT

<213> Homo sapiens

<400> 240

Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His 1 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50

<210> 241

<211> 58

<212> PRT

<213> Homo sapiens

<400> 241

Pro Glu Lys Asn Gly Ala Ala Thr Gly Met Asp Ala Ile Cys Ser His 1 10 15

Arg Leu Asp Pro Lys Ser Pro Gly Leu Asp Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr His Gly Ile Lys Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly 50 55

<210> 242

<211> 58

<212> PRT

<213> Homo sapiens

<400> 242

Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu 1 10 15

Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp Page 275 Glu Leu Ser Gln Leu Thr Asn Ser Val Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 243

<211> 58

<212> PRT

<213> Homo sapiens

<400> 243

Pro Glu Lys His Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Thr Leu
1 10 15

Arg Leu Asp Pro Thr Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 244

<211> 58

<212> PRT

<213> Homo sapiens

<400> 244

Pro Glu Lys His Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His 1 10 15

Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 245

<211> 58

<212> PRT

<213> Homo sapiens

<400> 245

Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His 1 10 15

Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asn Gly 50

<210> 246

<211> 58

<212> PRT

Homo sapiens

<400> 246

Pro Glu Lys Gln Glu Ala Ala Thr Gly Val Asp Thr Ile Cys Thr His 1 10 15

Arg Val Asp Pro Ile Gly Pro Gly Leu Asp Arg Glu Arg Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr Asn Ser Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asp Gly 50

<210> 247

58

<211> <212> **PRT**

<213> Homo sapiens

<400> 247

Pro Glu Lys Asp Lys Ala Ala Thr Arg Val Asp Ala Ile Cys Thr His 1 10 15

His Pro Asp Pro Gln Ser Pro Gly Leu Asn Arg Glu Gln Leu Tyr Trp 20 25 30

Glu Leu Ser Gln Leu Thr His Gly Ile Thr Glu Leu Gly Pro Tyr Thr 35 40 45

Leu Asp Arg Asp Ser Leu Tyr Val Asp Gly 50

```
<210> 248
<211>
       58
<212>
       PRT
```

<213> Homo sapiens

<400> 248

Ser Val Lys Asn Gly Ala Glu Thr Arg Val Asp Leu Leu Cys Thr Tyr 1 5 10 15

Leu Gln Pro Leu Ser Gly Pro Gly Leu Pro Ile Lys Gln Val Phe His 20 25 30

Glu Leu Ser Gln Gln Thr His Gly Ile Thr Arg Leu Gly Pro Tyr Ser 40 45

Leu Asp Lys Asp Ser Leu Tyr Leu Asn Gly 50

<210> 249

<211> <212> 58

PRT

<213> Homo sapiens

<400> 249

Pro Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Thr Thr Cys Thr Tyr 1 10 15

His Pro Asp Pro Val Gly Pro Gly Leu Asp Ile Gln Gln Leu Tyr Trp
20 25 30

Glu Leu Ser Gln Leu Thr His Gly Val Thr Gln Leu Gly Phe Tyr Val 35 40 45

Leu Asp Arg Asp Ser Leu Phe Ile Asn Gly

<210> 250

<211> 12

<212> PRT

<213> Homo sapiens

<400> 250

Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr $1 \hspace{1cm} 5 \hspace{1cm} 10$

<210> 251

<211> 12 <212> PRT

<213> Homo sapiens

<400> 251

```
Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Ile 1 	 5 	 10
<210> 252
<211> 12
<212> PRT
<213> Homo sapiens
<400> 252
Phe Thr His Arg Thr Ser Val Pro Thr Ser Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 253
<211> 12
<212> PRT
<213> Homo sapiens
<400> 253
Phe Thr His Arg Thr Ser Val Pro Thr Thr Ser Thr 10
<210> 254
<211>
<212>
        12
        PRT
<213>
         Homo sapiens
<400> 254
Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ser 1 10
<210> 255
<211> 12
<212> PRT
<213> Homo sapiens
<400> 255
Phe Thr His Arg Ser Ser Val Ser Thr Thr Ser Thr 1 5 10
<210> 256
<211> 12
<212> PRT
<213> Homo sapiens
<400> 256
Phe Thr His Arg Ser Ser Val Ala Pro Thr Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 257
<211> 12
         257
<212> PRT
```

```
09-965738substitute.ST25.txt
<213> Homo sapiens
<400> 257
Phe Thr His Arg Ser Ser Gly Leu Thr Thr Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 258
<211>
       12
<212>
        PRT
<213>
        Homo sapiens
<400> 258
Phe Thr His Arg Ser Phe Gly Leu Thr Thr Ser Thr 1 10
<210> 259
<211> 12
<212> PRT
<213> Homo sapiens
<400> 259
Phe Thr His Arg Ser Ser Phe Leu Thr Thr Ser Thr 1 \hspace{1cm} 10
<210> 260
<211> 12
<212> PRT
<213> Homo sapiens
<400> 260
Phe Thr His Arg Asn Phe Val Pro Ile Thr Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 261
<211> 12
<212> PRT
<213> Homo sapiens
<400> 261
Phe Thr His Arg Ser Ser Val Pro Thr Thr Ser Ile
1 5 10
<210>
        262
<211> 12
<212> PRT
<213> Homo sapiens
<400> 262
Phe Thr His Gln Ser Ser Val Ser Thr Thr Ser Thr 1 10
```

```
09-965738substitute.ST25.txt
<210> 263
<211>
       12
<212>
       PRT
<213> Homo sapiens
<400> 263
Phe Thr His Gln Thr Ser Ala Pro Asn Thr Ser Thr 1 10
<210> 264
<211> 12
<212> PRT
<213> Homo sapiens
<400> 264
Phe Thr His Gln Thr Phe Ala Pro Asn Thr Ser Thr 1 10
<210> 265
       12
<211>
<212> PRT
<213> Homo sapiens
<400> 265
Phe Thr His Gln Asn Ser Val Pro Thr Thr Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 266
<211> 12
<212> PRT
<213> Homo sapiens
<400> 266
Phe Thr His Gln Ser Ser Met Thr Thr Arg Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 267
<211>
       12
<212>
        PRT
<213> Homo sapiens
<400> 267
Phe Thr His Trp Ile Pro Val Pro Thr Ser Ser Thr 1 10
<210> 268
<211> 12
<212> PRT
<213> Homo sapiens
<400> 268
```

Phe Thr His Trp Ser Pro Ile Pro Thr Thr Ser Thr

Page 281

```
5
1
<210> 269
<211> 12
<212> PRT
<213> Homo sapiens
<400> 269
Phe Thr His Trp Ser Ser Gly Leu Thr Thr Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 270
<211> 12
<212> PRT
<213> Homo sapiens
<400> 270
Phe His Pro Arg Ser Ser Val Pro Thr Thr Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 271
<211> 12
<212> PRT
<213>
         Homo sapiens
<400> 271
Phe Asn Pro Arg Ser Ser Val Pro Thr Thr Ser Thr 1 10
<210> 272
<211> 12
<212> PRT
<213> Homo sapiens
<400> 272
Phe Asn Pro Trp Ser Ser Val Pro Thr Thr Ser Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 273
<211> 12
<212> PRT
<213> Homo sapiens
<400> 273
Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Ile
1 5 10
<210> 274
<211> 12
<212> PRT
<213> Homo sapiens
```

```
09-965738substitute.ST25.txt
<400> 274
Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Thr 1 10
<210> 275
<211> 12
<212> PRT
<213> Homo sapiens
<400> 275
Phe Thr Gln Arg Ser Ser Val Pro Thr Thr Ser Val 1 10
<210> 276
<211> 12
<212> PRT
<213> Homo sapiens
<400> 276
Tyr Asn Glu Pro Gly Leu Asp Glu Pro Pro Thr Thr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 277
<211> 12
<212> PRT
<213> Homo sapiens
<400> 277
Tyr Ala Pro Gln Asn Leu Ser Ile Arg Gly Glu Tyr 1 \hspace{1cm} 5 \hspace{1cm} 10
<210> 278
<211> 21
<212> PRT
<213> Homo sapiens
<400> 278
Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser 10 15
Ser Pro Ser Pro Thr
               20
<210> 279
<211> 23
<212> PRT
<213> Homo sapiens
<400> 279
```

Pro Gly Thr Ser Thr Val Asp Leu Arg Thr Ser Gly Thr Pro Ser Ser 10 15

Page 283

```
Leu Ser Ser Pro Thr Ile Met
              20
<210>
       280
<211>
        21
<212>
       PRT
<213>
       Homo sapiens
<400> 280
Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Phe Ser 1 10 15
Leu Pro Ser Pro Ala
              20
<210> 281
<211> 20
<212> PRT
<213> Homo sapiens
<400> 281
Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Ser Leu 1 \hspace{1cm} 10 \hspace{1cm} 15
Pro Ser Pro Thr
              20
<210> 282
<211> 20
<212> PRT
<213> Homo sapiens
<400> 282
Pro Gly Thr Ser Thr Val Asp Leu Gly Ser Gly Thr Pro Ser Leu Pro 1 \hspace{1cm} 10 \hspace{1cm} 15
Ser Ser Pro Thr
<210> 283
<211> 21
<212> PRT
<213> Homo sapiens
<400> 283
Pro Gly Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Ser 10 15
Leu Pro Ser Pro Thr
                                             Page 284
```

20

```
<210> 284
<211> 21
<212>
       PRT
<213> Homo sapiens
<400> 284
Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser 1 10 15
Lys Pro Gly Pro Ser
             20
<210> 285
<211> 21
<212> PRT
<213> Homo sapiens
<400> 285
Pro Trp Thr Ser Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Ser Pro 1 \hspace{1cm} 10 \hspace{1cm} 15
Val Pro Ser Pro Thr
<210> 286
<211> 21
<212> PRT
<213> Homo sapiens
<400> 286
Pro Gly Thr Ser Thr Val Tyr Trp Ala Thr Thr Gly Thr Pro Ser Ser 1 10 15
Phe Pro Gly His Thr
<210> 287
       21
<211>
<212>
       PRT
<213> Homo sapiens
<400> 287
Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Ser 1 10 15
Leu Pro Gly His Thr
20
```

```
09-965738substitute.ST25.txt
<210>
       288
<211>
       21
<212>
       PRT
<213>
       Homo sapiens
       288
<400>
Pro Gly Thr Ser Thr Val His Leu Ala Thr Ser Gly Thr Pro Ser Pro 1 10 15
Leu Pro Gly His Thr 20
<210>
       289
21
<211>
<212>
       PRT
<213> Homo sapiens
<400>
       289
Pro Asp Thr Ser Thr Met His Leu Ala Thr Ser Arg Thr Pro Ala Ser 1 10 15
Leu Ser Gly Pro Thr
<210> 290
<211> 21
<212> PRT
<213> Homo sapiens
<400>
       290
Pro Gly Thr Ser Ala Val His Leu Glu Thr Ser Gly Thr Pro Ala Ser 1 \hspace{1cm} 10 \hspace{1cm} 15
Leu Pro Gly His Thr 20
<210>
       291
<211>
       21
<212>
       PRT
<213>
       Homo sapiens
<400> 291
Pro Gly Thr Ser Ala Val His Leu Glu Thr Thr Gly Thr Pro Ser Ser 10 15
Phe Pro Gly His Thr
             20
<210>
        292
<211>
       21
<212> PRT
```

```
09-965738substitute.ST25.txt
<213> Homo sapiens
<400> 292
Pro Gly Thr Ser Thr Val His Leu Gly Thr Ser Glu Thr Pro Ser Ser 10 15
Leu Pro Arg Pro Ile
<210> 293
<211> 21
<212> PRT
<213> Homo sapiens
<400> 293
Pro Gly Thr Ser Ile Val Asn Leu Gly Thr Ser Gly Ile Pro Pro Ser 1 \hspace{1cm} 10 \hspace{1cm} 15
Leu Pro Glu Thr Thr
<210>
       294
<211> 21
<212> PRT
<213> Homo
       Homo sapiens
<400> 294
Pro Gly Thr Phe Thr Val Gln Pro Glu Thr Ser Glu Thr Pro Ser Ser 1 10 15
Leu Pro Gly Pro Thr 20
<210> 295
<211>
<212>
        21
        PRT
<213>
       Homo sapiens
<400> 295
Pro Gly Thr Pro Thr Val Asp Leu Gly Thr Ser Gly Thr Pro Val Ser 1 10 15
Lys Pro Gly Pro Ser
             20
<210> 296
<211>
       21
<212>
       PRT
<213>
      Homo sapiens
```

<400> 296

Pro Gly Thr Pro Thr Val Tyr Leu Gly Ala Ser Lys Thr Pro Ala Ser 1 10 15

Ile Phe Gly Pro Ser

297 <210>

<211> 16 <212> PRT

<213> Homo sapiens

<400> 297

Pro Lys Pro Ala Thr Thr Phe Leu Pro Pro Leu Ser Glu Ala Thr Thr 1 5 10 15

<210> 298 <211> 21

<212> PRT

<213> Homo sapiens

<400> 298

Gln Ile Asn Phe His Ile Val Asn Trp Asn Leu Ser Asn Pro Asp Pro 1 10 15

Thr Ser Ser Glu Tyr 20

<210>

<211> 1794 <212>

PRT <213> Homo sapiens

<400> 299

Met Glu His Ile Thr Lys Ile Pro Asn Glu Ala Ala His Arg Gly Thr 1 10 15

Ile Arg Pro Val Lys Gly Pro Gln Thr Ser Thr Ser Pro Ala Ser Pro 20 25 30

Lys Gly Leu His Thr Gly Gly Thr Lys Arg Met Glu Thr Thr Thr 35 40 45

Ala Leu Lys Thr Thr Thr Ala Leu Lys Thr Thr Ser Arg Ala Thr 50 60

Leu Thr Thr Ser Val Tyr Thr Pro Thr Leu Gly Thr Leu Thr Pro Leu 65 70 75 80

Asn Ala Ser Arg Gln Met Ala Ser Thr Ile Leu Thr Glu Met Met Ile Page 288

Thr Thr Pro Tyr Val Phe Pro Asp Val Pro Glu Thr Thr Ser Ser Leu 100 Ala Thr Ser Leu Gly Ala Glu Thr Ser Thr Ala Leu Pro Arg Thr Thr Pro Ser Val Leu Asn Arg Glu Ser Glu Thr Thr Ala Ser Leu Val Ser Arg Ser Gly Ala Glu Arg Ser Pro Val Ile Gln Thr Leu Asp Val Ser 145 150 155 160 Ser Ser Glu Pro Asp Thr Thr Ala Ser Trp Val Ile His Pro Ala Glu 165 170 175 Thr Ile Pro Thr Val Ser Lys Thr Thr Pro Asn Phe Phe His Ser Glu Leu Asp Thr Val Ser Ser Thr Ala Thr Ser His Gly Ala Asp Val Ser Ser Ala Ile Pro Thr Asn Ile Ser Pro Ser Glu Leu Asp Ala Leu Thr Pro Leu Val Thr Ile Ser Gly Thr Asp Thr Ser Thr Thr Phe Pro Thr 225 230 235 240 Leu Thr Lys Ser Pro His Glu Thr Glu Thr Arg Thr Thr Trp Leu Thr His Pro Ala Glu Thr Ser Ser Thr Ile Pro Arg Thr Ile Pro Asn Phe 260 265 270 260 Ser His His Glu Ser Asp Ala Thr Pro Ser Ile Ala Thr Ser Pro Gly Ala Glu Thr Ser Ser Ala Ile Pro Ile Met Thr Val Ser Pro Gly Ala 290 295 300 Glu Asp Leu Val Thr Ser Gln Val Thr Ser Ser Gly Thr Asp Arg Asn Met Thr Ile Pro Thr Leu Thr Leu Ser Pro Gly Glu Pro Lys Thr Ile

Ala	Ser	Leu	Val 340	Thr	His	Pro	09-9 Glu	96573 Ala 345	38sut Gln	Thr	ser Ser	ST25 Ser	Ala 350	Ile	Pro
Thr	Ser	Thr 355	Ile	Ser	Pro	Ala	Val 360	Ser	Arg	Leu	val	Thr 365	Ser	Met	val
Thr	Ser 370	Leu	Ala	Ala	Lys	Thr 375	Ser	Thr	Thr	Asn	Arg 380	Ala	Leu	Thr	Asn
Ser 385	Pro	Gly	Glu	Pro	Ala 390	Thr	Thr	val	Ser	Leu 395	val	Thr	His	Pro	Ala 400
Gln	Thr	Ser	Pro	Thr 405	√val	Pro	Тгр	Thr	Thr 410	Ser	Ile	Phe	Phe	ніs 415	Ser
Lys	Ser	Asp	Thr 420	Thr	Pro	Ser	Met	Thr 425	Thr	Ser	His	Gly	Ala 430	Glu	Ser
ser	Ser	Ala 435	val	Pro	Thr	Pro	Thr 440	val	Ser	Thr	Glu	Va1 445	Pro	Gly	val
Val	Thr 450	Pro	Leu	val	Thr	Ser 455	Ser	Arg	Ala	val	Ile 460	Ser	Thr	Thr	Ile
Pro 465	Ile	Leu	Thr	Leu	Ser 470	Pro	Gly	Glu	Pro	Glu 475	Thr	Thr	Pro	Ser	Met 480
Ala	Thr	Ser	His	Gly 485	Glu	Glu	Ala	Ser	ser 490	Ala	Ile	Pro	Thr	Pro 495	Thr
∨al	ser	Pro	Gly 500	val	Pro	Gly	٧al	va1 505	Thr	Ser	Leu	val	Thr 510	Ser	Ser
Arg	Аlа	Val 515	Thr	ser	Thr	Thr	11e 520	Pro	Ile	Leu	Thr	Phe 525	Ser	Leu	Gly
Glu	Pro 530	Glu	Thr	Thr	Pro	Ser 535	Met	Аlа	Thr	Ser	ніs 540	Gly	Thr	Glu	Ala
Gly 545	Ser	Ala	val	Pro	Thr 550	∨al	Leu	Pro	Glu	Val 555	Pro	Gly	Met	val	Thr 560
Ser	Leu	val	Ala	ser 565	Ser	Arg	Ala	val	Thr 570	ser	Thr	Thr	Leu	Pro 575	Thr
Leu	Thr	Leu	Ser 580	Pro	Gly	Glu	Pro	G]u 585	Thr	Thr	Pro	Ser	Met 590	Ala	Thr

Ser His Gly Ala Glu Ala Ser Ser Thr Val Pro Thr Val Ser Pro Glu 600 Val Pro Gly Val Val Thr Ser Leu Val Thr Ser Ser Gly Val Asn Ser Thr Ser Ile Pro Thr Leu Ile Leu Ser Pro Gly Glu Leu Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Ala Glu Ala Ser Ser Ala Val Pro Thr Pro Thr Val Ser Pro Gly Val Ser Gly Val Val Thr Pro Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr Thr Ile Pro Ile Leu Thr 680 Leu Ser Ser Ser Glu Pro Glu Thr Thr Pro Ser Met Ala Thr Ser His Gly Val Glu Ala Ser Ser Ala Val Leu Thr Val Ser Pro Glu Val Pro Gly Met Val Thr Ser Leu Val Thr Ser Ser Arg Ala Val Thr Ser Thr 725 730 735 Thr Ile Pro Thr Leu Thr Ile Ser Ser Asp Glu Pro Glu Thr Thr Ser Leu Val Thr His Ser Glu Ala Lys Met Ile Ser Ala Ile Pro Thr 755 760 765 Leu Ala Val Ser Pro Thr Val Gln Gly Leu Val Thr Ser Leu Val Thr Ser Ser Gly Ser Glu Thr Ser Ala Phe Ser Asn Leu Thr Val Ala Ser Ser Gln Pro Glu Thr Ile Asp Ser Trp Val Ala His Pro Gly Thr Glu Ala Ser Ser Val Val Pro Thr Leu Thr Val Ser Thr Gly Glu Pro Phe 820 Thr Asn Ile Ser Leu Val Thr His Pro Ala Glu Ser Ser Ser Thr Leu 840

Page 291

- Pro Arg Thr Thr Ser Arg Phe Ser His Ser Glu Leu Asp Thr Met Pro 850 855 860
- Ser Thr Val Thr Ser Pro Glu Ala Glu Ser Ser Ser Ala Ile Ser Thr 865 870 875 880
- Thr Ile Ser Pro Gly Ile Pro Gly Val Leu Thr Ser Leu Val Thr Ser 885 890 895
- Ser Gly Arg Asp Ile Ser Ala Thr Phe Pro Thr Val Pro Glu Ser Pro 900 905 910
- His Glu Ser Glu Ala Thr Ala Ser Trp Val Thr His Pro Ala Val Thr 915 920 925
- Ser Thr Thr Val Pro Arg Thr Thr Pro Asn Tyr Ser His Ser Glu Pro 930 935 940
- Asp Thr Thr Pro Ser Ile Ala Thr Ser Pro Gly Ala Glu Ala Thr Ser 945 955 960
- Asp Phe Pro Thr Ile Thr Val Ser Pro Asp Val Pro Asp Met Val Thr 965 970 975
- Ser Gln Val Thr Ser Ser Gly Thr Asp Thr Ser Ile Thr Ile Pro Thr 980 985 990
- Leu Thr Leu Ser Ser Gly Glu Pro Glu Thr Thr Ser Phe Ile Thr 995 1000 1005
- Tyr Ser Glu Thr His Thr Ser Ser Ala Ile Pro Thr Leu Pro Val 1010 1015 1020
- Ser Pro Gly Ala Ser Lys Met Leu Thr Ser Leu Val Ile Ser Ser 1025 1030 1035
- Gly Thr Asp Ser Thr Thr Thr Phe Pro Thr Leu Thr Glu Thr Pro 1040 1045 1050
- Tyr Glu Pro Glu Thr Thr Ala Ile Gln Leu Ile His Pro Ala Glu 1055 1060 1065
- Thr Asn Thr Met Val Pro Arg Thr Thr Pro Lys Phe Ser His Ser 1070 1075 1080
- Lys Ser Asp Thr Thr Leu Pro Val Ala Ile Thr Ser Pro Gly Pro Page 292

Glu Ala Ser Ser Ala Val Ser Thr Thr Thr Ile Ser Pro Asp Met 1105 1100 Ser Asp Leu Val Thr Ser Leu Val Pro Ser Ser Gly Thr Asp Thr 1115 1120 1125 Ser Thr Thr Phe Pro Thr Leu Ser Glu Thr Pro Tyr Glu Pro Glu Thr Thr Ala Thr Trp Leu Thr His Pro Ala Glu Thr Ser Thr Thr 1145 1150 1155 val Ser Gly Thr Ile Pro Asn Phe Ser His Arg Gly Ser Asp Thr Ala Pro Ser Met Val Thr Ser Pro Gly Val Asp Thr Arg Ser Gly Val Pro Thr Thr Thr Ile Pro Pro Ser Ile Pro Gly Val Val Thr Ser Gln Val Thr Ser Ser Ala Thr Asp Thr Ser Thr Ala Ile Pro 1215 1205 Thr Leu Thr Pro Ser Pro Gly Glu Pro Glu Thr Thr Ala Ser Ser 1220 1225 1230 Ala Thr His Pro Gly Thr Gln Thr Gly Phe Thr Val Pro Ile Arg 1235 Thr Val Pro Ser Ser Glu Pro Asp Thr Met Ala Ser Trp Val Thr 1250 His Pro Pro Gln Thr Ser Thr Pro Val Ser Arg Thr Thr Ser Ser 1270 Phe Ser His Ser Ser Pro Asp Ala Thr Pro Val Met Ala Thr Ser 1280 1285 1290 1280 Pro Arg Thr Glu Ala Ser Ser Ala Val Leu Thr Thr Ile Ser Pro

Gly Ala Pro Glu Met Val Thr Ser Gln Ile Thr Ser Ser Gly Ala

1315

09-965738substitute.ST25.txt Ala Thr Ser Thr Thr Val Pro Thr Leu Thr His Ser Pro Gly Met Pro Glu Thr Thr Ala Leu Leu Ser Thr His Pro Arg Thr Glu Thr Ser Lys Thr Phe Pro Ala Ser Thr Val Phe Pro Gln Val Ser Glu Thr Thr Ala Ser Leu Thr Ile Arg Pro Gly Ala Glu Thr Ser Thr Ala Leu Pro Thr Gln Thr Thr Ser Ser Leu Phe Thr Leu Leu Val 1390 1395 1385 Thr Gly Thr Ser Arg Val Asp Leu Ser Pro Thr Ala Ser Pro Gly 1400 Val Ser Ala Lys Thr Ala Pro Leu Ser Thr His Pro Gly Thr Glu 1415 1420 Thr Ser Thr Met Ile Pro Thr Ser Thr Leu Ser Leu Gly Leu Leu 1435 Glu Thr Thr Gly Leu Leu Ala Thr Ser Ser Ser Ala Glu Thr Ser 1445 1450 1455 Thr Ser Thr Leu Thr Leu Thr Val Ser Pro Ala Val Ser Gly Leu Ser Ser Ala Ser Ile Thr Thr Asp Lys Pro Gln Thr Val Thr Ser Trp Asn Thr Glu Thr Ser Pro Ser Val Thr Ser Val Gly Pro Pro Glu Phe Ser Arg Thr Val Thr Gly Thr Thr Met Thr Leu Ile Pro 1515 Ser Glu Met Pro Thr Pro Pro Lys Thr Ser His Gly Glu Gly Val 1525 1520 Ser Pro Thr Thr Ile Leu Arg Thr Thr Met Val Glu Ala Thr Asn 1535 1540 1545 Leu Ala Thr Thr Gly Ser Ser Pro Thr Val Ala Lys Thr Thr Thr 1550 1560 1550 Page 294

Thr Phe Asn Thr Leu Ala Gly Ser Leu Phe Thr Pro Leu Thr Thr 1575 Pro Gly Met Ser Thr Leu Ala Ser Glu Ser Val Thr Ser Arg Thr 1580 1585 1590 Ser Tyr Asn His Arg Ser Trp $\,$ Ile Ser Thr Thr Ser $\,$ Ser Tyr Asn $\,$ 1595 $\,$ 1600 $\,$ 1605 Arg Arg Tyr Trp Thr Pro Ala Thr Ser Thr Pro Val Thr Ser Thr Phe Ser Pro Gly Ile Ser Thr Ser Ser Ile Pro Ser Ser Thr Ala 1630 Ala Thr Val Pro Phe Met Val Pro Phe Thr Leu Asn Phe Thr Ile Thr Asn Leu Gln Tyr Glu Glu Asp Met Arg His Pro Gly Ser Arg Lys Phe Asn Ala Thr Glu Arg Glu Leu Gln Gly Leu Leu Lys Pro 1670 1680 Leu Phe Arg Asn Ser Ser Leu Glu Tyr Leu Tyr Ser Gly Cys Arg 1685 1690 1695 1690 Leu Ala Ser Leu Arg Pro Glu Lys Asp Ser Ser Ala Met Ala Val 1700 Asp Ala Ile Cys Thr His Arg Pro Asp Pro Glu Asp Leu Gly Leu 1715 1720 1725 Asp Arg Glu Arg Leu Tyr Trp Glu Leu Ser Asn Leu Thr Asn Gly 1730 1740 Ile Gln Glu Leu Gly Pro Tyr Thr Leu Asp Arg Asn Ser Leu Tyr 1745 1750 1755 Val Asn Gly Phe Thr His Arg Ser Ser Met Pro Thr Thr Ser Thr 1760 1765 1770Pro Gly Thr Ser Thr Val Asp Val Gly Thr Ser Gly Thr Pro Ser Ser Ser Pro Ser Pro Thr

1790

300 <210> <211> 284 <212> **PRT** Homo sapiens <400> 300 Ile Thr Leu Leu Arg Asp Ile Gln Asp Lys Val Thr Thr Leu Tyr Lys
1 10 15 Gly Ser Gln Leu His Asp Thr Phe Arg Phe Cys Leu Val Thr Asn Leu 20 25 30 Thr Met Asp Ser Val Leu Val Thr Val Lys Ala Leu Phe Ser Ser Asn 35 40 45 Leu Asp Pro Ser Leu Val Glu Gln Val Phe Leu Asp Lys Thr Leu Asn 50 55 60 Ala Ser Phe His Trp Leu Gly Ser Thr Tyr Gln Leu Val Asp Ile His 65 70 75 80 Val Thr Glu Met Glu Ser Ser Val Tyr Gln Pro Thr Ser Ser Ser Ser 90 95 Thr Gln His Phe Tyr Leu Asn Phe Thr Ile Thr Asn Leu Pro Tyr Ser 100 105 110 Gln Asp Lys Ala Gln Pro Gly Thr Thr Asn Tyr Gln Arg Asn Lys Arg 115 120 125 Asn Ile Glu Asp Ala Leu Asn Gln Leu Phe Arg Asn Ser Ser Ile Lys Ser Tyr Phe Ser Asp Cys Gln Val Ser Thr Phe Arg Ser Val Pro Asn 145 150 155 160 Arg His His Thr Gly Val Asp Ser Leu Cys Asn Phe Ser Pro Leu Ala 165 170 175 Arg Arg Val Asp Arg Val Ala Ile Tyr Glu Glu Phe Leu Arg Met Thr 180 185 190 Arg Asn Gly Thr Gln Leu Gln Asn Phe Thr Leu Asp Arg Ser Ser Val 195 200 205 Leu Val Asp Gly Tyr Ser Pro Asn Arg Asn Glu Pro Leu Thr Gly Asn Page 296

Ser As 225	p Leu Pro	Phe	Trp 230	Ala	val	Ile	Leu	Ile 235	Glу	Leu	Ala	Gly	Leu 240		
Leu Gl	y Leu Ile	Thr 245	Cys	Leu	Ile	Cys	Gly 250	٧a٦	Leu	val	Thr	Thr 255	Arg		
Arg Ar	g Lys Lys 260		Gly	Glu	Tyr	Asn 265	٧a٦	Gln	Gln	Gln	Cys 270	Pro	Gly		
Tyr Ty	r Gln Ser 275	His	Leu	Asp	Leu 280	Glu	Asp	Leu	Gln						
<210> <211> <212> <213>	301 24 DNA Artifici	al So	equei	nce											
<220> <223> Synthetic primer															
<400> 301 gtctctatgt caatggtttc accc													24		
<210> 302 <211> 21 <212> DNA <213> Artificial Sequence															
<220> <223> Synthetic primer															
<400> 302 tagctgctct ctgtccagtc c													21		
<210> 303 <211> 22 <212> DNA <213> Artificial Sequence															
<220> <223>	Syntheti	c pr	imer												
<400> 303 ggacaaggtc accacactct ac												22			
<210> <211> <212> <213>	304 24 DNA Artifici	al S	eque	nce											
<220> <223>															
<400>	0> 304 Page 297														

09-965738substitute.ST25.txt gcagatcctc caggtctagg tgtg	24
<210> 305 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> Synthetic primer	
<400> 305 gtctctatgt caatggtttc accc	24
<210> 306 <211> 21 <212> DNA <213> Artificial Sequence	
<220> <223> Synthetic primer	
<400> 306 tagctgctct ctgtccagtc c	21
<210> 307 <211> 468 <212> DNA <213> Homo sapiens	
<400> 307 actgctggcc ctctcctggt gccattcacc ctcaacttca ccatcaccaa cctgcagtat	60
gaggaggaca tgcatcgccc tggatctagg aagttcaaca ccacagagag ggtcctgcag	120
ggtctgctta gtcccatatt caagaacacc agtgttggcc ctctgtactc tggctgcaga	180
ctgacctctc tcaggtctga gaaggatgga gcagccactg gagtggatgc catctgcatc	240
catcatcttg accccaaaag ccctggactc aacagagagc ggcțgtactg ggagctgagc	300
cgactgacca atggcatcaa agagctgggc ccctacaccc tggacaggaa cagtctctat	360
gtcaatggtt tcacccatcg gacctctgtg cccaccacca gcactcctgg gacctccaca	420
gtggaccttg gaacctcagg gactccattc tccctcccaa gccccgca	468
<210> 308 <211> 156 <212> PRT <213> Homo sapiens	
<400> 308	
Thr Ala Gly Pro Leu Leu Val Pro Phe Thr Leu Asn Phe Thr Ile Thr 1 15	
Asn Leu Gln Tyr Glu Glu Asp Met His Arg Pro Gly Ser Arg Lys Phe 20 25 30	

Page 298

Asn Thr Thr Glu Arg Val Leu Gln Gly Leu Leu Ser Pro Ile Phe Lys 35 40 45

Asn Thr Ser Val Gly Pro Leu Tyr Ser Gly Cys Arg Leu Thr Ser Leu 50 60

Arg Ser Glu Lys Asp Gly Ala Ala Thr Gly Val Asp Ala Ile Cys Ile 65 70 75 80

His His Leu Asp Pro Lys Ser Pro Gly Leu Asn Arg Glu Arg Leu Tyr 85 90 95

Trp Glu Leu Ser Arg Leu Thr Asn Gly Ile Lys Glu Leu Gly Pro Tyr 100 105 110

Thr Leu Asp Arg Asn Ser Leu Tyr Val Asn Gly Phe Thr His Arg Thr 115 120 125

Ser Val Pro Thr Thr Ser Thr Pro Gly Thr Ser Thr Val Asp Leu Gly 130 140

Thr Ser Gly Thr Pro Phe Ser Leu Pro Ser Pro Ala 145 150 155

<210> 309

<211> 10

<212> PRT

<213> Homo sapiens

<400> 309

Pro Gly Ser Arg Lys Phe Lys Thr Thr Glu 1 5 10

<210> 310

<211> 5

<212> DNA

<213> Homo sapiens

<400> 310

aataa

5